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by *Philippe Nimmegeers, Joost Lauwers, Dries Telen, Filip Logist and Jan Van Impe*

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Identifiability of large-scale non-linear dynamic network models applied to the ADM1-case study

Philippe Nimmegeers^a, Joost Lauwers^a, Dries Telen^a, Filip Logist^a, Jan Van Impe^{a,b}

^a*KU Leuven, Department of Chemical Engineering, BioTeC+ & OPTEC, Gebroeders De Smetstraat 1, 9000 Ghent, Belgium*

^b*Corresponding author. jan.vanimpe@kuleuven.be
Tel: +32.16.32.14.66*

Abstract

In this work, both the structural and practical identifiability of the Anaerobic Digestion Model no. 1 (ADM1) is investigated, which serves as a relevant case study of large non-linear dynamic network models. The structural identifiability is investigated using the *probabilistic* algorithm, adapted to deal with the specifics of the case study (i.e., a large-scale non-linear dynamic system of differential and algebraic equations). The practical identifiability is analyzed using a Monte Carlo parameter estimation procedure for a ‘non-informative’ and ‘informative’ experiment, which are heuristically designed.

The model structure of ADM1 has been modified by replacing parameters by parameter combinations, to provide a generally locally structurally identifiable version of ADM1. This means that in an idealized theoretical situation, the parameters can be estimated accurately. Furthermore, the generally positive structural identifiability results can be explained from the large number of interconnections between the states in the network structure. This interconnectivity, however, is also observed in the parameter estimates, making uncorrelated parameter estimations in practice difficult.

Keywords: Structural identifiability, Practical identifiability, Anaerobic Digestion Model No.1, Non-linear biological models

1. Introduction

- Dynamic models are essential tools for the study of large-scale bioprocesses. Due to the complexity of the processes, however, these models generally contain large numbers of (i) non-linear ordinary differential or differential alge-

5 braic equations (ODEs or DAEs) and (ii) parameters which have to be esti-
6 mated. The latter issue raises questions whether the parameters can be esti-
7 mated univocally (i.e., unique parameter estimates) and with sufficient accu-
8 racy and precision (which is reflected in the width of the confidence intervals
9 on the parameter estimates). Furthermore the parameter estimates should be
10 consistent with respect to parameter values that are reported in literature.

11 The problem of non-uniqueness of parameter estimates is a question of *identi-*
12 *fiability*. This concept acts on two levels: the *structural identifiability* and the
13 *practical identifiability*. The former refers to the structure of the model which
14 does or does not allow the unique characterization of the parameters, even un-
15 der perfect ‘identification conditions’. A lack of structural identifiability could
16 be attributed to a poor choice of model structure such as overparameteriza-
17 tion or the inclusion of phenomena that can never be experimentally verified
18 or refuted. The practical identifiability refers to finding accurate and precise
19 parameter estimates considering the influence of issues such as the choice and
20 limitation of the applied input and the measuring conditions. Since the condi-
21 tions for structural identifiability are stricter than for practical identifiability,
22 structural identifiability is in fact a *necessary condition* for practical identifi-
23 bility. For a model that is not structurally identifiable, it is impossible to find
24 reliable parameter estimates.

25 In this work the Anaerobic Digestion Model No. 1 (ADM1) [1] is considered
26 as a case study of a large-scale non-linear biological network model. For the
27 structural identifiability analysis: different measurement and input scenarios
28 are evaluated. For the practical identifiability, the single situation in which
29 all states are measured and all possible inputs are available, i.e., the situation
30 that yields the best estimates, is investigated.

31 The contributions of this article are three-fold. Firstly, to show how an anal-
32 ysis of the structural and practical identifiability of a large-scale model con-
33 sisting of DAEs and non-rational expressions can be performed. Secondly, to
34 yield structural and practical identifiability results for ADM1, which are to
35 our knowledge not available. Finally, to generalize the results of the ADM1
36 case study to dynamic network models: to investigate the structural aspects
37 of the model causing the (non-)identifiability.

38 The outline of the work is as follows: first the ADM1 case study is presented.
39 This also includes a statement of the identifiability problem and an overview
40 of the analyzed cases concerning the applied inputs and measured outputs. In
41 the next sections definitions concerning structural identifiability are stated, to-
42 gether with a description of the method to study it. This is followed by a de-
43 scription of the followed methodology for analyzing the practical identifiabil-
44 ity. The results section starts with an overview of the structural identifiabil-
45 ity results for the ADM1 case study. The structural aspects of the case study
46 causing the (non-)identifiability are highlighted. Next, the results concerning
47 the practical identifiability are presented, with a highlight on the structural
48 aspects of the model causing the (non-)identifiability. The conclusions are pre-
49 sented in the last section.

50 2. Case study

51 Anaerobic Digestion Model No. 1 (ADM1) is a generic model developed by
52 the International Water Association's Anaerobic Digestion Modelling Task
53 Group [2, 1]. The ADM1 model knows numerous simulation applications in
54 literature varying from anaerobic digestion of organic wastes to waste wa-

ter treatment (e.g., [3, 4, 5, 6]). Therefore it can be seen as the state-of-the-art model in anaerobic digestion. The ADM1 model network structure is depicted in Figure ??, based on the implementation of the equations presented in Appendix A [7]. The model comprises 39 states: 24 soluble components (S_i , with $i \in \{\text{su, aa, fa, va, bu, pro, ac, h2, ch4, IC, IN, I}\}$), 3 gaseous components ($S_{\text{gas},i}$, with $i \in \{\text{h2, ch4, co2}\}$), 5 particulate composites (X_i , with $i \in \{\text{xc, ch, pr, li, I}\}$) with i being a component and 7 microbial groups (X_j , with $j \in \{\text{su, aa, fa, c4, pro, ac, h2}\}$), degrading component j . Since the reactions related to the acid-base equilibrium can be considered instantaneous, they are expressed in algebraic expressions (AEs) of index 1, following [1]. Most of the states are expressed either in terms of Chemical Oxygen Demand (COD) or kmol C or N per m^3 . See Table 1 of Appendix A for an overview of all states.

ADM1 includes 22 stoichiometric, 19 composition and 37 kinetic parameters. The stoichiometric parameters express the ratios in which COD is converted from one state into another: (i) 7 biomass on component yield coefficients ($Y_i \in [0, 1]$, with i being the component consumed by the microbial group) and (ii) 15 component on component yield coefficients, ($f_{i,j} \in [0, 1]$ with i being the component produced from component j). The composition parameters N_i and C_i express the nitrogen and carbon content of a component i . They act as conversion factors between COD-based states and the inorganic nitrogen and carbon. Of the 19 composition parameters, 5 are considered to be known, i.e., the carbon content of CH_4 , valeric acid, butyric acid, propionic acid and acetic acid. The kinetic parameters appear in the reaction rates of the model. They include 11 first-order constants, 7 Monod maximum growth rate and 7 saturation constants, 5 parameters concerning secondary substrate uptake or inhibition, 3 pairs of upper and lower pH-boundaries which appear

82 in the pH-inhibition expressions and 1 liquid-gas mass transfer parameter. See
83 Table 3 to 7 of Additional file 1 for an overview of all parameters.

84 In addition, there are several physico-chemical constants such as the acid-base
85 dissociation constants and operational parameters, i.e., the liquid volume and
86 the temperature. It is assumed that they are known and they have the default
87 value listed in Table 7 and 8 in Appendix B [7].

88 Note, that in this work, more parameters are considered ‘unknown’ than typi-
89 cally the case in anaerobic digestion applications [8, 9]. The structural identi-
90 fiability in these applications, however, can be deduced from the investigated
91 more general situation. If a parameter is identifiable in the latter situation,
92 it will also be identifiable when part of the set of parameters are assigned a
93 numerical value.

94 Concerning the inputs, there are two types: the liquid inflow q_{liq} and the con-
95 centrations of the components in the inflow. Both the liquid inflow as the con-
96 centrations of the inflow are controlled. The numerical values used for the in-
97 puts are listed in Table 3 of Additional File 2 [7], except for $X_{su,in}$ which is
98 changed from 0.0 to 0.1.

99 Concerning the output, it is for now assumed that all states in ADM1, both
100 differential and algebraic, can be measured. This is in accordance with the
101 underlying philosophy of this manuscript in finding an *upper limit* on the
102 number of parameters that can be estimated and the corresponding quality of
103 the estimates. For the practical identifiability analysis, the situation yielding
104 the most informative option is chosen: both the dilution rate and the liquid
105 flow rate are controlled and all states are measured.

106 *Remark:* In practice, liquid inflow, gaseous outflow and the dilution rate can
107 be measured. Typically, liquid samples are taken daily, e.g. [10],[11], and some
108 state values have to be deduced from measurements such as total COD and
109 alkalinity. An interesting direction of research from a practitioner's point of
110 view, could be to relax these assumptions and analyze the identifiability when
111 only typical measurements of a digester are performed: biogas production and
112 composition, pH, total and soluble chemical oxygen demand (COD), volatile
113 fatty acids (VFA), total organic carbon, total inorganic carbon, Kjeldahl-
114 Nitrogen, total inorganic nitrogen, total phosphorous, orthophosphate, to-
115 tal alkalinity, total solids, total volatile solids. Using the measurement-to-
116 state mapping of Kleerebezem and Van Loosdrecht (2006) [12] or Zaher et
117 al. (2009) [13] these measurements can be converted into the ADM1 states.
118 A difficulty here is that direct measurement of the distinct microbial groups
119 is not possible and including it requires estimation of the initial conditions
120 of the different microbial groups. However, major developments in absolute
121 quantification using quantitative polymerase chain reaction (qPCR) technique
122 and fluorescence in situ hybridization (FISH) could create opportunities for
123 measurement of these groups. See for instance Junicke et al. (2014) [14].

124 3. Mathematical methods

125 In the following section the concepts of structural and practical identifiabil-
126 ity are introduced. In addition, several different methods to investigate these
127 concepts are presented.

3.1. Structural identifiability

Structural identifiability is defined following the definitions of [15]. Consider the dynamic model Σ of the form:

$$\Sigma = \begin{cases} \frac{d\mathbf{x}}{dt} = \mathbf{f}(\mathbf{x}, \mathbf{z}, \boldsymbol{\theta}, \mathbf{u}, t) & (1) \\ \mathbf{0} = \mathbf{g}(\mathbf{x}, \mathbf{z}, \boldsymbol{\theta}, \mathbf{u}, t) & (2) \\ \mathbf{y} = \mathbf{h}(\mathbf{x}, \mathbf{z}, \boldsymbol{\theta}, \mathbf{u}, t) & (3) \\ \mathbf{x}(0) = \mathbf{x}_0 & (4) \end{cases}$$

where $\mathbf{x} = [x_1, \dots, x_{n_x}]^T \in \mathbb{R}^{n_x}$ the differential state vector, $\mathbf{z} = [z_1, \dots, z_{n_z}]^T \in \mathbb{R}^{n_z}$ the algebraic state vector $\boldsymbol{\theta} = [\theta_1, \dots, \theta_{n_\theta}]^T \in \mathbb{R}^{n_\theta}$ the parameter vector, $\mathbf{u} = [u_1, \dots, u_{n_u}]^T \in \mathbb{R}^{n_u}$ the input vector, $\mathbf{y} = [y_1, \dots, y_{n_y}]^T \in \mathbb{R}^{n_y}$ the output vector, $t \in \mathbb{R}$ the time, \mathbf{f} a \mathbb{R}^{n_x} vector function, \mathbf{g} a \mathbb{R}^{n_z} vector function, \mathbf{h} a \mathbb{R}^{n_y} vector function expressing the output and $\mathbf{x}_0 \in \mathbb{R}^{n_x}$ the initial state vector. It is assumed that for each system described by (1)-(4) there exists a set of true parameter values $\boldsymbol{\theta}^*$, which are considered to be unknown. Parameter estimation is then equivalent to finding parameter estimates $\hat{\boldsymbol{\theta}}$, that are as close as possible to the true parameter values $\boldsymbol{\theta}^*$.

Structural identifiability then relates to the question whether the true parameter values can be found from the input-output behavior, denoted with $M(\boldsymbol{\theta})$. In studying this relationship, the following idealized conditions are assumed:

- (i) the model is a perfect representation of the process, (ii) all data is noise-free and (iii) the input \mathbf{u} can be chosen at will and (iv) the sampling is continuous. Formally, a parameter θ_{i_θ} , $i_\theta = \{1, \dots, n_\theta\}$ is *structurally globally identifiable* if under these ideal conditions and for almost any true parameter value $\boldsymbol{\theta}^*$, the true parameter values can be uniquely estimated from input-

148 output behavior M , i.e.,

$$M(\hat{\theta}) = M(\theta^*) \quad (5)$$

149 has the unique solution:

$$\hat{\theta}_{i_\theta} = \theta_{i_\theta}^*. \quad (6)$$

150 If all parameters of a model Σ are structurally globally identifiable, the model
 151 Σ is structurally globally identifiable. With ‘for almost any θ^* ’ it is meant
 152 that structural identifiability may vary depending on the true parameter val-
 153 ues. Often, this will only occur in singular points of the feasible parameter
 154 space. A parameter θ_{i_θ} is said to be *structurally locally identifiable* if (5) has a
 155 countable number of solutions (larger than 1) [16]. A model Σ is structurally
 156 locally identifiable if all its parameters are structurally locally identifiable. A
 157 parameter θ_{i_θ} , $i_\theta = \{1, \dots, n_\theta\}$ is *structurally unidentifiable* or *non-identifiable*
 158 if under the previously presented idealized conditions and for almost any true
 159 parameter value θ^* , (5) has an infinite number of solutions. A model Σ is
 160 structurally unidentifiable if one of its parameters is structurally unidentifi-
 161 able. If a model is unidentifiable, it is often possible to find structurally iden-
 162 tifiable parameter combinations.

163 Different techniques exist to determine the structural identifiability of dy-
 164 namic models, e.g., investigating the coefficients of the Taylor expansion [17]
 165 or investigating generators of the differential ideal described by the model
 166 equations [18]. In general, these techniques work well for small-scale systems,
 167 yielding a general result for the structural identifiability. However, due to the
 168 extensive symbolic calculations involved, they do not deal well with large sys-
 169 tems, mainly due to memory considerations [19]. Therefore, the procedure
 170 followed in this work is based on the *probabilistic* algorithm of Sedoglavic [20],
 171 since it has been proven to be applicable for several large-scale models [21].

172 The method elaborates on reformulating the identifiability problem as a prob-
 173 lem of observability by considering the parameters as non-varying states and
 174 checking whether the initial conditions $[\mathbf{x}_0^T, \boldsymbol{\theta}^T]^T$ are *locally algebraically ob-*
 175 *servable*. Without going into the details, it can be shown [22, 23, 24] that lo-
 176 cal algebraic observability is a sufficient condition for structural local identifia-
 177 bility. Other names used in literature for local algebraic observability are *local*
 178 *distinguishability* [22], Hermann-Krener observability or *HK-observability* [25],
 179 *local generic observability* [23] and the Exact Arithmetic Rank (EAR) condi-
 180 tion [26]. For brevity, this condition will be referred to as *observability* for the
 181 remainder of the text.

182 In what follows the method used to investigate the observability of $[\mathbf{x}_0^T, \boldsymbol{\theta}^T]^T$ is
 183 briefly presented. This is followed by several modifications of the model and
 184 method needed in order to analyze the structural identifiability of ADM1. Fi-
 185 nally, the procedure for identifying parameter combinations is explained.

186 3.1.1. Methodology

187 The probabilistic method applies to ODE systems, i.e., $n_z = 0$ for which \mathbf{h}
 188 and \mathbf{f} are rational functions in the parameters, states and inputs. It is as-
 189 sumed that the initial conditions \mathbf{x}_0 do not depend on the parameters. The
 190 initial conditions \mathbf{x}_0 and parameters $\boldsymbol{\theta}$ of the system Σ are observable if the
 191 rank of the following Jacobian evaluated for $[\mathbf{x}_0^{*T}, \boldsymbol{\theta}^{*T}]$ equals the total num-

ber of parameters and states, i.e., $n_x + n_\theta$:

$$\begin{bmatrix} \frac{\partial h_1}{\partial x_1} & \dots & \frac{\partial h_1}{\partial x_{n_x}} & \frac{\partial h_1}{\partial \theta_1} & \dots & \frac{\partial h_1}{\partial \theta_{n_\theta}} \\ \vdots & \vdots & \vdots & \vdots & \vdots & \vdots \\ \frac{\partial h_{n_y}}{\partial x_1} & \dots & \frac{\partial h_{n_y}}{\partial x_{n_x}} & \frac{\partial h_{n_y}}{\partial \theta_1} & \dots & \frac{\partial h_{n_y}}{\partial \theta_{n_\theta}} \\ \frac{\partial \mathcal{L}_{\mathbf{f}} h_1}{\partial x_1} & \dots & \frac{\partial \mathcal{L}_{\mathbf{f}} h_1}{\partial x_{n_x}} & \frac{\partial \mathcal{L}_{\mathbf{f}} h_1}{\partial \theta_1} & \dots & \frac{\partial \mathcal{L}_{\mathbf{f}} h_1}{\partial \theta_{n_\theta}} \\ \vdots & \vdots & \vdots & \vdots & \vdots & \vdots \\ \frac{\partial \mathcal{L}_{\mathbf{f}} h_{n_y}}{\partial x_1} & \dots & \frac{\partial \mathcal{L}_{\mathbf{f}} h_{n_y}}{\partial x_{n_x}} & \frac{\partial \mathcal{L}_{\mathbf{f}} h_{n_y}}{\partial \theta_1} & \dots & \frac{\partial \mathcal{L}_{\mathbf{f}} h_{n_y}}{\partial \theta_{n_\theta}} \\ \vdots & \vdots & \vdots & \vdots & \vdots & \vdots \\ \frac{\partial \mathcal{L}_{\mathbf{f}}^{(n_x+n_\theta-1)} h_1}{\partial x_1} & \dots & \frac{\partial \mathcal{L}_{\mathbf{f}}^{(n_x+n_\theta-1)} h_1}{\partial x_{n_x}} & \frac{\partial \mathcal{L}_{\mathbf{f}}^{(n_x+n_\theta-1)} h_1}{\partial \theta_1} & \dots & \frac{\partial \mathcal{L}_{\mathbf{f}}^{(n_x+n_\theta-1)} h_1}{\partial \theta_{n_\theta}} \\ \vdots & \vdots & \vdots & \vdots & \vdots & \vdots \\ \frac{\partial \mathcal{L}_{\mathbf{f}}^{(n_x+n_\theta-1)} h_{n_y}}{\partial x_1} & \dots & \frac{\partial \mathcal{L}_{\mathbf{f}}^{(n_x+n_\theta-1)} h_{n_y}}{\partial x_{n_x}} & \frac{\partial \mathcal{L}_{\mathbf{f}}^{(n_x+n_\theta-1)} h_{n_y}}{\partial \theta_1} & \dots & \frac{\partial \mathcal{L}_{\mathbf{f}}^{(n_x+n_\theta-1)} h_{n_y}}{\partial \theta_{n_\theta}} \end{bmatrix} \quad (7)$$

in which the Lie-derivative on the vector field \mathbf{f} is introduced:

$$\mathcal{L}_{\mathbf{f}} := \frac{\partial}{\partial t} + \sum_{i=1}^{n_x} f_i \frac{\partial}{\partial x_i} + \sum_{j \in \mathbb{N}} \sum_{i=1}^{n_u} u_i^{(j+1)} \frac{\partial}{\partial u_i^{(j)}} \quad (8)$$

with:

$$\mathcal{L}_{\mathbf{f}}^{(j)} = \underbrace{\mathcal{L}_{\mathbf{f}} \mathcal{L}_{\mathbf{f}} \dots \mathcal{L}_{\mathbf{f}}}_{j \text{ times}} \quad (9)$$

and $u_i^{(j)}$ the j -th derivative of u_i . To determine if a specific state/parameter is observable, it suffices to eliminate the corresponding column in (7) and see whether the rank of (7) changes.

A fully symbolic calculation of the Jacobian matrix is theoretically possible, but the equations quickly become huge. The algorithm of [20] calculates the elements of the matrix in a numerical way using rationals.

First, integers are assigned to the parameters and initial conditions. Next, the input is specified to a random integer coefficient power series. Using this input, the specified initial conditions and parameter values, a truncated power

series of the states, as well as their sensitivity to the initial conditions and parameters, are calculated. Using these power series sensitivities, values for the Lie-derivatives are calculated [20]. To prevent calculations with large integer numbers, all calculations are done modulo a large prime number.

It should be remarked that the result is a *sufficient* condition for structural local identifiability. The method does not guarantee that a parameter is structurally globally identifiable. These caveats make that the method gives less general results than for instance other structural identifiability techniques, e.g. differential algebra [27].

3.1.2. Implementation aspects

Concerning the implementation of the method for the case of ADM1, three aspects are addressed:

1. The relations between the stoichiometric parameters to ascertain a correct COD balance in the system, have to be taken into account (Equations (10) to (12)). Therefore $f_{pr,xc}$, $f_{h2,aa}$ and $f_{h2,su}$ by the expressions in terms of the other stoichiometric parameters.

$$1 = f_{xI,xc} + f_{sI,xc} + f_{ch,xc} + f_{pr,xc} + f_{li,xc} \quad (10)$$

$$1 = f_{pro,su} + f_{ac,su} + f_{h2,su} + f_{bu,su} \quad (11)$$

$$1 = f_{va,aa} + f_{bu,aa} + f_{pro,aa} + f_{ac,aa} + f_{h2,aa} \quad (12)$$

2. ADM1 consists of AEs which in principle cannot be analyzed with the adopted method. Therefore, instead of the 10 index-1 algebraic equa-

222 tions, their underlying ODEs are used, i.e.,

$$\mathbf{0} = \mathbf{g}(\mathbf{x}, \boldsymbol{\theta}, \mathbf{z}, \mathbf{u}, t) \quad (13)$$

223 is converted into

$$\frac{d\mathbf{z}}{dt} = - \left(\frac{\partial \mathbf{g}}{\partial \mathbf{z}} \right)^{-1} \left(\frac{\partial \mathbf{g}}{\partial \mathbf{x}} \right) \left(\frac{\partial \mathbf{x}}{\partial t} \right) = - \left(\frac{\partial \mathbf{g}}{\partial \mathbf{z}} \right)^{-1} \left(\frac{\partial \mathbf{g}}{\partial \mathbf{x}} \right) \mathbf{f} = \mathbf{f}^\dagger(\mathbf{x}, \boldsymbol{\theta}, \mathbf{z}, \mathbf{u}, t). \quad (14)$$

224 The initial conditions of these differentiated algebraic states are deter-
 225 mined by solving the algebraic equations, evaluated with the initial con-
 226 ditions of the differential states in Appendix B [7]. All values are con-
 227 verted to an exact rational representation.

228 3. The method is not able to deal with non-rational expressions although
 229 these appear in ADM1. Therefore the non-rational expressions for the
 230 pH-inhibition factors $I_{\text{pH},i}$ are reformulated by introducing the param-
 231 eters $k_{\text{pH},i}$ and the auxiliary state ω_i , $i = \{1, \dots, 3\}$:

$$I_{\text{pH},i} = \frac{K_{\text{pH},i}^{n_{\text{pH},i}}}{S_{h^+}^{n_{\text{pH},i}} + K_{\text{pH},i}^{n_{\text{pH},i}}} \rightarrow I_{\text{pH},i} = \frac{k_{\text{pH},i}}{\omega_i + k_{\text{pH},i}}, \quad k_{\text{pH},i} = K_{\text{pH},i}^{n_{\text{pH},i}}, \quad \omega_i = S_{h^+}^{n_{\text{pH},i}}. \quad (15)$$

232 The auxiliary state ω_i is added to the equations of the system:

$$\frac{d\omega_i}{dt} = n_{\text{pH},i} \frac{\omega_i}{S_{h^+}} \cdot \frac{dS_{h^+}}{dt}. \quad (16)$$

233 Concerning the identifiability analysis, instead of the 3 pairs of upper
 234 and lower limits, the 3 pairs of parameters $n_{\text{pH},i}$ and $k_{\text{pH},i}$ are now in-
 235 vestigated. However, it has to be stressed that in none of the scenarios,
 236 the 3 states ω_i are measured. Analogously as for the differentiated alge-
 237 braic states, the initial conditions of the ω_i and the parameter values are

numerically assigned to comply with (15).

Remark: The aforementioned adaptations impose implicit relations between the initial conditions and parameter values. which may lead to conflicting results. For instance, the method may assign ω_i and $k_{\text{pH},i}$ to be non-observable while $S_{\text{h}+}$ and $n_{\text{pH},i}$ are found to be observable. Clearly, from the observability of $S_{\text{h}+}$ and $n_{\text{pH},i}$ and (15) ω_i is also observable. Using $k_{\text{pH},i} = K_{\text{pH},i}^{n_{\text{pH},i}}$ and $\omega_i = S_{\text{h}+}^{n_{\text{pH},i}}$, it can then be deduced that $\text{pH}_{\text{UL},i} - \text{pH}_{\text{LL},i}$ is observable. The observability analysis is then repeated, with the observable parameters assigned a numerical value, in order to detect other observable parameters. The results presented are those after solving such *conflicting* results.

If the system is found to non-observable, one could try to find observable parameter combinations. Non-observability occurs when points on a direction in the parametric space render the input-output invariant, i.e., a so-called *symmetry*. It can be shown (see [22, 20]) that these symmetries can be calculated by calculating the kernel of the Jacobian matrix of which the columns associated with observable states/parameters are omitted.

In case the number of non-observable parameters/states is low, it is often much easier just to check whether the proposed parameter/state combinations involving non-observable parameters/states are observable, by considering their combination as a single parameter. These parameter combinations can often be discovered after close examination of the model equations. This approach is followed in this work.

260 3.2. Practical identifiability

261 Structural identifiability is a necessary but not sufficient condition for finding
262 univocal, accurate and precise parameter estimates. Experimental artefacts
263 such as the nature and magnitude of the noise, the choice of applied inputs
264 and measurements, the sampling rate and the form of the input may confound
265 the solution, i.e., increase the uncertainty of the estimate. In this work, the
266 practical identifiability analysis assesses the effect of the type of applied input
267 and the magnitude of the noise levels.

268 The practical identifiability is assessed by performing parameter estimations
269 on *in silico* data, i.e., data generated by simulations of the model.

270 In what follows a short description of parameter estimations and the proce-
271 dure on how the quality of the estimates is assessed, are presented. This is
272 followed by a description on how the experiments are designed.

273 3.2.1. Parameter estimation and quantification of uncertainty

274 The parameter estimations are based on the maximum likelihood criterion
275 which searches the value of the parameter vector θ that gives the highest like-
276 lihood to the observed data, considering the error distribution on the mea-
277 surements. Under the assumptions of (i) no characterization error, (ii) in-
278 dependently normally distributed measurements noise and (iii) known and
279 univariate variance, it can be proven that under these assumptions the max-
280 imization of the log-likelihood function (the logarithm of the likelihood func-
281 tion) corresponds to the minimization given in Equation (17) [15]. Note that
282 for other distributions different maximum likelihood estimators would be ob-

283 tained.

$$\min_{\boldsymbol{\theta}} \frac{1}{2} \sum_{i_y=1}^{n_y} \sum_{i=1}^{n_t} \frac{1}{\sigma_{i,i_y}^2} \left(y_{i_y}^s(t_i) - y_{i_y}(t_i, \boldsymbol{\theta}) \right)^2 \quad (17)$$

284 with n_t the number of time points t_i , σ_{i,i_y}^2 the measurement variance on the
 285 measured values $y_{i_y}^s(t_i)$ and $y_{i_y}(t_i, \boldsymbol{\theta})$ the predicted values.

286 In order to quantify the quality of the estimates, Monte Carlo resampling
 287 analyses are performed. The rationale is that a set of measurements \mathbf{y}^s is a
 288 particular realization of a *true* profile on which random noise is superimposed.
 289 The practical identifiability is then based on the estimates of the finite num-
 290 ber of realizations of \mathbf{y}^s . Practically, an *in silico* reference profile is generated
 291 using the true parameter value $\boldsymbol{\theta}^*$.

292 Note that the inverse of the Fisher Information Matrix (FIM) defined by the
 293 model and the experimental design could also be used as a measure for the
 294 quality of the estimates. However, the Fisher information matrix only pro-
 295 vides a linear approximation, while Monte Carlo simulations provide a real
 296 measure of the actual distribution of the parameter estimates in a setting with
 297 measurement noise. Furthermore, the authors opted for Monte Carlo simula-
 298 tions over the Fisher information matrix computation as it involves only the
 299 repetitions of multiple parameter estimations while for the FIM also the sensi-
 300 tivities with respect to the parameters are required. For the first experiment,
 301 the FIM approach would result in 2900 (i.e. number of estimated parameters
 302 (71 parameters+29 initial conditions)×29 dynamic states) in addition to the
 303 original states.

304 The following statistics are used: the mean $\bar{\boldsymbol{\theta}}_{MC}$, the standard deviation $\boldsymbol{\sigma}_{MC}$

and the covariance matrix \mathbf{P}_{MC} of the n_{it} realizations:

$$\bar{\theta}_{\text{MC}} = \sum_{i=1}^{n_{\text{it}}} \frac{1}{n_{\text{it}}} \hat{\theta}_i \quad (18)$$

$$\sigma_{\text{MC}} = \sqrt{\sum_{i=1}^{n_{\text{it}}} \frac{1}{n_{\text{it}}} (\hat{\theta}_i - \bar{\theta}_{\text{MC}})^2} \quad (19)$$

$$\mathbf{P}_{\text{MC}} = \sum_{i=1}^{n_{\text{it}}} (\hat{\theta}_i - \bar{\theta}_{\text{MC}})(\hat{\theta}_i - \bar{\theta}_{\text{MC}}). \quad (20)$$

These are then used to calculate the relative bias $\text{Bias}_{\text{rel}, i_\theta}$ and the relative standard deviation $\sigma_{\text{rel}, i_\theta}$ for each parameter or initial condition i_θ :

$$\text{Bias}_{\text{rel}, i_\theta} = \frac{\theta_{i_\theta}^* - \bar{\theta}_{\text{MC}, i_\theta}}{\theta_{i_\theta}^*} \quad (21)$$

$$\sigma_{\text{rel}, i_\theta} = \frac{\sigma_{\text{MC}, i_\theta}}{\theta_{i_\theta}^*}. \quad (22)$$

For large-scale models with many parameters, inspection of the different elements of the covariance matrix can be difficult to interpret. Alternatively, the matrix \mathbf{P}'_{MC} with elements $P'_{\text{MC}, i_\theta, j_\theta}$, $i_\theta = 1, \dots, n_\theta + n_x$, $j_\theta = 1, \dots, n_\theta + n_x$ which consists of roots of the absolute value of the normalized covariances, is calculated:

$$P'_{\text{MC}, i_\theta, j_\theta} = \sqrt{\left| \frac{P_{\text{MC}, i_\theta, j_\theta}}{\theta_{i_\theta}^* \theta_{j_\theta}^*} \right|}. \quad (23)$$

In this way, the diagonal elements of \mathbf{P}'_{MC} correspond with the absolute value of the relative standard deviations. Note that this measure looks very similar to the Pearson correlation coefficient ($R_{\text{Pearson}} = \frac{P_{\text{MC}, i_\theta, j_\theta}}{\sigma_{i_\theta} \sigma_{j_\theta}}$, with σ_{i_θ} and σ_{j_θ} the standard deviations of the i_θ -th and j_θ -th parameter), a measure for the linear dependence between different variables. Since ADM1 is a highly

nonlinear model and a nonlinear dependence between the parameters has to be covered, the authors considered normalized covariances to investigate the (nonlinear) dependence between different variables.

The number of realizations was chosen such that the addition of an extra realization changes the Monte Carlo estimates $\bar{\theta}_{MC,i_\theta}$ and σ_{MC,i_θ} less than 2%. This is mostly reached at around 100 iterations.

The overall quality of the Monte Carlo analysis is based on the average normalized root mean squared errors, R_1 , the normalized average error between the predictions and the measurements and R_2 , the normalized average error between the predictions and the true values. To assess the simulated measurements, the average normalized mean squared error between the measured and the true values R_3 is calculated:

$$R_1 = \sqrt{\frac{1}{n_{it}} \frac{1}{n_t} \frac{1}{n_y} \sum_{i=1}^{n_{it}} \sum_{t=1}^{n_t} \sum_{y=1}^{n_y} \frac{(y_{i_y,i_t} - y_{i_y,i_t}^s)^2}{\sigma_{i_y}^2}} \quad (24)$$

$$R_2 = \sqrt{\frac{1}{n_{it}} \frac{1}{n_t} \frac{1}{n_y} \sum_{i=1}^{n_{it}} \sum_{t=1}^{n_t} \sum_{y=1}^{n_y} \frac{(y_{i_y,i_t} - y_{i_y,i_t}^*)^2}{\sigma_{i_y}^2}} \quad (25)$$

$$R_3 = \sqrt{\frac{1}{n_{it}} \frac{1}{n_t} \frac{1}{n_y} \sum_{i=1}^{n_{it}} \sum_{t=1}^{n_t} \sum_{y=1}^{n_y} \frac{(y_{i_y,i_t}^s - y_{i_y,i_t}^*)^2}{\sigma_{i_y}^2}} \quad (26)$$

3.2.2. Design of experiments

The experiment is designed in order to find the ‘best’ profile to estimate the parameters, concerning the choice of outputs, inputs and the applied input profile within the experimental boundaries, e.g., maximum concentrations,

339 duration of experiment. In that way, an upper limit concerning the practi-
 340 cal identifiability is sought, i.e., to show that the parameter *can* be uniquely
 341 identified on experimental data. The design is made on heuristic rules and
 342 guidelines.

343 The heuristic rules focus on obtaining accurate estimates for the kinetic pa-
 344 rameters. Primarily because these parameters appear in a non-linear way in
 345 the equations, and are thus, are the most challenging in the estimation.

346 The general guideline followed is that the concentrations of the states involved
 347 in the reaction kinetics, including inhibitory functions, are such that the com-
 348 plete range of different kinetic values are active.

349 More specifically, for Monod-type kinetics, shown in Equation (27), the adop-
 350 tion of this general guideline is logical.

$$\mu(S) = \frac{\mu_{max}S}{K_S + S} \quad (27)$$

351 For very small concentrations of substrate S , $\rho \approx \mu_{max}/K_S$. For high con-
 352 centrations $\rho \approx \mu_{max}$. Thus, in order to estimate both parameters well, both
 353 extremes have to be reached approximately. Several sources [28, 29, 30] rec-
 354 ommend $S < 0.1K_S$ and $S > K_S$ or $S > 2K_S$.

355 For the secondary substrate uptake and inhibition functions, the value of the
 356 associated uptake or inhibition constant will only have an effect if the sec-
 357 ondary substrate concentrations are low and high respectively. For the prac-
 358 tical identifiability of these functions, the requirement is that the secondary

359 substrate has low concentrations for the secondary substrate uptake functions
360 or high concentrations for the secondary substrate inhibition functions.

361 Practically, the evaluation of an experiment concerning the practical identifi-
362 ability of the kinetic parameters will be determined by plotting the range of
363 concentrations of the states on the kinetic expressions.

364 3.3. Software

365 The structural identifiability tests are performed using the Maple `observabilityTest`
366 tool of [20], with the aforementioned adjustments to take into account the im-
367 plementation of AEs and non-rational expressions.

368 The parameter estimation problem is a dynamic optimization problem, which
369 is solved in an in-house developed software called Pomodoro [31], [32]. This
370 software can be downloaded from [http://cit.kuleuven.be/biotec/software/](http://cit.kuleuven.be/biotec/software/pomodoro)
371 `pomodoro` or <https://perswww.kuleuven.be/~u0093798/software.php> and
372 the cited work [31], can be found on [http://www.student.kuleuven.be/](http://www.student.kuleuven.be/~s0212066/pomodoro/)
373 `~s0212066/pomodoro/`.

374 States and controls are discretized following an orthogonal collocation frame-
375 work. In this approach, the value of the states are, like the parameters and
376 the initial conditions, considered to be optimization variables [33]. The con-
377 trol variables are discretized as piecewise constant. This usually results
378 in a very large but structured non-linear programming (NLP) problem. The
379 maximum number of allowed iterations was 2000 and the relative convergence
380 tolerance was $1E-7$.

381 To increase calculation speed and stability, all states and parameters in the

parameter estimation loop have been normalized. Additionally, a small value of $1\text{E}-10$ is added to the denominator ($S_{\text{bu}} + S_{\text{va}}$) in Equation (8) and (9) of Appendix A to avoid division by zero due to numerical errors. To prevent too sudden changes of input, changes in input variables are ‘spread’ in a ramp over the course of 1 day (i.e., the derivative of the input with respect to time is constrained).

The assigned lower and upper bounds for the optimization variables are the following:

- parameters: $\theta^*/4$ and $4 \times \theta^*$, with an upper limit of 0.999 for the stoichiometric parameters
- states: $0.2 \times \min \mathbf{x}^*(t)$ and $1.8 \times \max \mathbf{x}^*(t)$, $t = 0, \dots, n_t$
- initial conditions: $\mathbf{x}(0)^*/2$ and $\mathbf{x}(0)^* \times 2$

with θ^* the true parameter value and $\mathbf{x}(t)^*$ the true profile, i.e., without noise. As initial guess, $\mathbf{x}(t)^*$ was given for the states and initial conditions and θ^* for the parameters. Additionally, the relationships between the stoichiometric parameters of Equations (10) to (12) are given as extra constraints on the parameters. Finally, the differences between the lower and upper pH-inhibition levels, i.e., $\text{pH}_{\text{UL,aa}} - \text{pH}_{\text{LL,aa}}$, $\text{pH}_{\text{UL,h2}} - \text{pH}_{\text{LL,h2}}$ and $\text{pH}_{\text{UL,ac}} - \text{pH}_{\text{LL,ac}}$, are constrained to be less than 2.5.

All computations are done on a Intel® Core™ i7 – 3770, with 8 CPUs at 3.40 GHz each and 16 Gb of RAM.

4. Results and discussion

4.1. Structural identifiability

The results of the observability analyses of the different scenarios are summarized in Table 1 and discussed next. Each observability analysis of ADM1 took between 35-45 minutes to calculate.

Table 1: The results of the parameters of ADM1 for the scenarios with inputs q_{liq} and/or $S_{x,in}$. Observability is equivalent to structurally locally identifiable.

Measurement	Result
All states	All parameters observable excl. C_{sI} , C_{xI}
Any state excl. S_I , X_I , S_{cat+} , S_{an-}	All parameters observable excl. $f_{sI,xc}$, $f_{xI,xc}$, C_{sI} , C_{xI} , $S_I(0)$, $X_I(0)$, $S_{cat+}(0)$, $S_{an-}(0)$
S_I	All parameters observable excl. C_{sI} , C_{xI} , $X_I(0)$, $S_{cat+}(0)$, $S_{an-}(0)$
X_I	All parameters observable excl. C_{sI} , C_{xI} , $S_I(0)$, $S_{cat+}(0)$, $S_{an-}(0)$
S_{cat+}	None observable excl. $S_{cat+}(0)$
S_{an-}	None observable excl. $S_{an-}(0)$

Concerning the observability results, it is found that in all six measurement cases in Table 1, the non-observable parameters are related to the soluble and particulate inerts (i.e., the carbon compositions C_{sI} and C_{xI} and the stoichiometric yield of COD from particulate components to soluble $f_{sI,xc}$ and particulate inerts $f_{xI,xc}$). Since only the total free inorganic carbon is measured in the system, it is impossible to determine how much carbon is fixed in either the particulate or soluble inerts. Thus, the non-observability of the two parameters results from the structure of the model. In addition, if particulate or soluble inerts are not measured, it cannot be determined how much COD is fixed in each of the states, since they are both dead ends in the degradation pathway (??). This explains the non-observability of $f_{xI,xc}$ and $f_{sI,xc}$.

In the cases in which the corresponding states are not measured, the initial concentrations of soluble inerts (S_I), particulate inerts (X_I), cations (S_{cat+}) and anions (S_{an-}) are not observable. Furthermore, if only cation or anion concentrations are measured, no parameters are observable due to the fact that these states *only* ‘interact’ with the other states in the ion-balance of Equation (87) of Appendix A. In this equation, only their difference $S_{cat+} - S_{an-}$ appears. Hence, without measuring the anions or cations separately, these states cannot be estimated separately.

4.1.1. Parameter combinations

Because the results in Table 1 show several non-observable parameters/states, it is of interest to look for observable parameter combinations that involve the non-observable parameters/states. Although mathematical techniques exist to identify these parameter combinations automatically, i.e., the calculation of Lie point symmetries (see [20]), the limited number of non-observable parameters/states enables to manually check parameter/state combinations such that there is no need to apply them in the case of ADM1. The proposed parameter combinations are listed in Table 2 and their observability is verified by substituting the mentioned non-observable parameters by the accompanying parameter combination and recalculating the observability of the system. The original model structure and the structure with the two first proposed parameter combinations are shown in Figure 2. Note that no combination of states exists to estimate $S_I(0)$ or $X_I(0)$ when these states are not measured.

Table 2: Observable parameter combinations, based on the results of Table 1. Note that observability is equivalent to structurally locally identifiable.

Non-observable parameters/states	Observable parameter/state combination
C_{sI}, C_{xI}	$(f_{sI,xc}C_{sI} + f_{xI,xc}C_{xI})$
$f_{sI,xc}, f_{xI,xc}$	$(f_{sI,xc} + f_{xI,xc})$
S_{cat+}, S_{an-}	$(S_{an-} - S_{cat+})$

4.1.2. Structural aspects

The found results can be explained from the model structure.

Firstly, as already mentioned, the properties of the states that are dead ends in the model structure are not observable, unless they are measured directly. Secondly, the good structural identifiability properties can be attributed to the large number of interactions between the different states: forward *and* backward (i.e., for reversible reaction fluxes). These interactions are discussed in depth in Appendix C.

4.2. Practical identifiability

As discussed in the Methods section, the influence of the choice of the applied input and the magnitude of noise are investigated. The first aspect will be handled by investigation of a 'non-informative' and an 'informative' experiment. For each of the two experiments different noise levels are analyzed. Since it was found that the parameters C_I and C_{xI} are not structurally identifiable, the structurally identifiable parameter combination $C_I = f_{sI,xc}C_{sI} + f_{xI,xc}C_{xI}$ is estimated instead. In addition, the role of the N-limiting parameter $K_{IN,lim}$ is solely numerical and it will not be included in the parameter estimation procedure. Hence, 71 parameters and 29 initial conditions (i.e., corresponding to the 29 original dynamic states) that need to be estimated.

460 The outline of these sections is as follows: First, the general experimental and
 461 process conditions are stated. This is followed by an overview of the designed
 462 experiments. Next, the results are presented and discussed. Finally, the re-
 463 sults for ADM1 are generalized.

464 4.2.1. *Experimental and process conditions*

465 In order to work with a structurally (locally) identifiable model, the parame-
 466 ter C_I as defined in Table 2 is used instead of the composition parameters C_{sI}
 467 and C_{xI} .

468 The initial conditions are the obtained steady-states when the liquid inflow
 469 as described in Appendix B is applied with a volumetric flow rate $q_{liq} = 85$
 470 m^3d^{-1} . This relatively low rate induces lower biomass concentrations such
 471 that the system is easier to ‘excite’, i.e., induce a change in concentrations.
 472 This steady-state is calculated by simulation over a period of 1000 days.

473 The noise is normally distributed $\mathcal{N}(0, \sigma_{i_y}^2)$ with a constant variance for each
 474 output y_{i_y} . The variance of the distribution equals the square of a set per-
 475 centage of the initial conditions. Different percentage levels will be investi-
 476 gated.

477 Samples are taken 4 times per day, which is a high number for an anaerobic
 478 digestion system. Higher values are not considered due to the additional com-
 479 putational burden.

480 The parameter $K_{IN,lim}$ associated with N as a limiting substrate is assigned
 481 its true parameter value of $1E-4$, since this parameter is more of a numerical

482 nature than of a physical one [7].

483 For this experiment design, all state variables are considered to be measured.
 484 Admittedly, this is unrealistic but helps establishing an upper limit on what
 485 can be estimated for ADM1.

486 Concerning the applied input of the system, several boundaries are set. The
 487 applied controls are the liquid inflow q_{liq} , the input concentrations of cations
 488 $S_{\text{cat}^+, \text{in}}$ and $S_{\text{an}^-, \text{in}}$ representing added bases and acids, monosaccharides
 489 $S_{\text{su}, \text{in}}$ and amino acids $S_{\text{aa}, \text{in}}$. It will be seen that the control of just these
 490 variables, leads to satisfactory results concerning the specified design speci-
 491 fications. Based on the solubility of monosaccharides and amino acids, a max-
 492 imum concentration for $S_{\text{su}, \text{in}}$ and $S_{\text{aa}, \text{in}}$ is determined. In that respect, the
 493 monosaccharides are assumed to be D-glucose with a maximum solubility at
 494 35°C of 612 gCOD L⁻¹ [34] and the amino acids are assumed to be a mixture
 495 of amino acids originating from the decomposition of casein with a maximum
 496 solubility of about 20 gCOD L⁻¹. Cations and anions are added in the form
 497 concentrated strong bases and acids, e.g., NaOH and HCl such as reported by
 498 [35]. There are no specific limitations on the total amount of added substrate.

499 4.2.2. Overview of performed experiments

500 Based on the discussed conditions and limitations, two experiments are de-
 501 signed. The first acts as a non-informative experiment, in which only the dilu-
 502 tion rate (or hydraulic retention time) is changed.

503 *Experiment 1.*

504 The motivation for this first experiment is to get an idea on what is achiev-

able when performing only a simple control action. At the beginning of this experiment, the liquid inflow q_{liq} is doubled from $85 \text{ m}^3\text{d}^{-1}$ to $170 \text{ m}^3\text{d}^{-1}$ so that the dilution rate d increases from 0.025 d^{-1} to 0.05 d^{-1} . This dilution rate is maintained for 50 days so that the system evolves from the steady-state SS_1 to steady-state SS_2 . This is depicted in Figure 3. Note that for this experiment 100 parameters are estimated (i.e., 71 parameters and 29 initial conditions).

Experiment 2.

In the second experiment, the heuristic guidelines discussed in Section 3.2.2 are applied. In order to activate the full spectrum of the kinetic expressions, a second experiment is proposed in which a change in liquid inflow is combined with the addition of extra intermediate components in the liquid feed while taking into account the previously considered experimental limitations. To prevent the extra inflow of intermediate components to be directly consumed by the available biomass and leading to accompanying biomass growth, the reaction rates are kept low by reducing the pH to 4 by adding acid in the form of anions (S_{an^-}). At this pH, the substrate uptake rates are practically zero. The concentration of monosaccharides and amino acids are then raised, taking into consideration the limits as determined in Section 4.2.1. This is maintained for 10 days. Afterwards, the pH is then gradually raised to the starting point by adding bases in the form of cations (S_{cat^+}), as well as the liquid inflow q_{liq} which is changed from $85 \text{ m}^3\text{d}^{-1}$ to $170 \text{ m}^3\text{d}^{-1}$ after 25 days. The applied input profiles for $S_{\text{su,in}}$, $S_{\text{aa,in}}$, q_{liq} and the desired pH profile are shown in Figure 4. Note that for this experiment 98 parameters are estimated (i.e., 71 parameters and 27 initial conditions).

For both experiments, four different noise levels are imposed. The standard

deviations of the Gaussian noise are 5%, 10%, 15% and 25% of the initial values listed in Table 2 of Appendix B.

A successful parameter estimation took between 5 and 20 minutes.

Several numerical issues did arise during the implementation of this specific experiment. This concerns the tracking of the desired pH-profile and the concentrations of inorganic carbon and hydrogen. Following modifications have been applied to the model to address these issues:

1. To enable the desired pH-profile, the concentration profiles of S_{an^-} , in - S_{cat^+} , in depicted in Figure 5 has to be applied. Since the required changes in this profile are too sudden to apply the desired pH-profile in a piecewise fashion with time step of 0.25 d, the pH-profile $\text{pH}(t)$ itself is enforced in the model. Furthermore, the algebraic equation concerning the ion balance in Equation (87) of Appendix A has been changed to:

$$0 = 100 \cdot [S_{\text{h}^+}(t) - 10^{-\text{pH}(t)}]. \quad (28)$$

The factor 100 is inserted for a better tracking. Conceptually, this kind of alterations can be thought of as the introduction of a *perfect* controller, which instantaneously feeds the acids and bases to the system, according to Figure 5. Note that due to this tracking, the states S_{an^-} and S_{cat^+} become irrelevant and are removed from the equations.

2. A limiting substrate inhibition for H_2 (S_{h_2}) and inorganic carbon (S_{IC}) are included to prevent negative concentrations for S_{h_2} and S_{IC} . Note that such functions were already included for inorganic nitrogen S_{IN} in

the form of $K_{IN,lim}$. The used inhibition functions are the following:

$$I_{IC,lim} = \frac{S_{IC}}{S_{IC} + K_{IC,lim}} \quad (29)$$

$$I_{h2,lim} = \frac{S_{h2}}{S_{h2} + K_{h2,lim}} \quad (30)$$

with $K_{h2,lim} = 10^{-5}$ and $K_{h2,lim} = 10^{-8}$. These factors are added to the S_{h2} and S_{IC} consuming reactions. These are the uptake of lipids S_{li} in ρ_4 , the uptake of valeric acid S_{vva} in ρ_8 , the uptake of butyric acid S_{bu} in ρ_9 and the uptake of hydrogen S_{h2} for S_{IC} . This is also the single S_{h2} consuming reaction. The inclusion of these inhibition factors has no effect on the state profiles for experiment 1.

The majority of the resulting state profiles are depicted, together with a more in-depth discussion in Appendix D. The resulting state profiles are plotted on the kinetic expressions in Figure 6. It is clear that for all kinetic expressions, the values of the states are spread out over the entire spectrum. Importantly, for the Monod type kinetics, the concentrations lie both above and below the reference concentration of the saturation constant, which corresponds with a rate of $\mu_{max}/2$.

Parameter estimation results

In this section the parameter estimation results for the two experiments at the 4 noise levels are presented and discussed. A Monte Carlo parameter estimate will be labeled ‘good’, ‘mediocre’ or ‘bad’ when its relative bias or standard deviation, as defined in Equation (21) and (22) is below 5%, between 5% and

572 25% and above 25%, respectively.

573 *Experiment 1.*

574 The performance of the Monte Carlo analysis for the 4 different noise levels,
 575 i.e., 5%, 10%, 15% and 25% is presented in Table 3 with the statistics defined
 576 in Equations (24) to (25). Since $R_1 \approx 1$, the error between the measurement
 577 and the predictions is almost equivalent to measurement error. The error be-
 578 tween the true and predicted values are about 10 %, which is acceptable to
 579 good.

Table 3: Performance of the Monte Carlo analysis for experiment 1.

Noise	R_1	R_2	R_3
5%	0.9945	0.1002	0.9996
10%	0.9878	0.0848	0.9926
15%	0.9895	0.0841	0.9937
25%	0.9901	0.0902	0.9929

580 The parameter estimation results are listed in Table 1 and 2 of Appendix E in
 581 which the bias and the observed standard deviation are reported. Generally
 582 speaking, the results are rather poor, and as expected, the parameter esti-
 583 mates become worse with increasing measurement noise.

584 In order to asses the independence of the parameter estimates, the ‘relative
 585 covariance’ \mathbf{P}'_{MC} as defined Equation (23) is constructed. For brevity, only the
 586 5% noise level case is calculated, depicted in Figure 7. There are 2720 pairs
 587 with no or weak interdependence (less than 5 %), 1746 pairs with moderate
 588 interdependence (between 5 and 25%)and 484 pairs of strongly related (more
 589 than 25%) parameter estimates in the upper triangle of the matrix. This does
 590 not include the diagonal elements of the matrix which represent absolute val-
 591 ues of the single parameter relative standard deviations. The lower triangle is
 592 the mirror image of the upper triangle.

593 There are several trends that can be observed. First, there is generally a con-
 594 siderable interdependence between the parameters. Also, the parameters that
 595 were found to have large relative standard deviations, are also the ones that
 596 have strong relations with others. In Figure 7 it can be seen that if a row or
 597 column has dark marker on the diagonal, it is very likely to have another dark
 598 or gray marker on its row or column.

599 There are two blocks of parameters with little interdependence to other pa-
 600 rameters. This can be seen on the figure as columns or rows with only a rather
 601 limited amount of gray markers. These are the first order kinetic parameters
 602 (36.-40.) and the initial conditions (72.-100.). Additionally, there are several
 603 blocks of parameters that generally have intermediate interdependence with
 604 other parameters. These are the pH-inhibition levels (66.-71.), the LCFA from
 605 lipids yield $f_{fa,li}$ (1.) and the biomass yield coefficients (2.-8.).

606 The rest of the parameter estimates generally show substantial interdepen-
 607 dence, although several big blocks of particularly strong interdependence can
 608 be observed. These consist of the relations between the kinetic parameters
 609 concerning S_{su} , S_{aa} , S_{pro} , S_{fa} , S_{bu} , S_{va} , S_{h2} , S_{ac} (41.-58.) , the associated sto-
 610 ichiometric parameters (14.-22.) and all composition parameters. Exceptions
 611 are the saturation constant for the acetic acid degradation $K_{S,ac}$ (55.) and
 612 the carbon content of the inerts C_I (28.). In Appendix E, these results are
 613 explained from the structure and the equations.

614 In conclusion, this simple experiment showed several problems concerning the
 615 practical identifiability of the parameters of ADM1. Because of the rather lim-
 616 ited spread of the states on the kinetic expressions, most kinetic parameters
 617 could not be estimated properly. Moreover, many of the parameter estimates

show significant interdependence between each other. Nevertheless, for this simple experiment, adequate results were obtained for the disintegration kinetics, the hydrolysis kinetics, the liquid-gas mass transfer constant and the initial conditions of the states. Also, the saturation constant for acetate uptake is reasonably well estimated. It is safe to say that, if these parameters can be estimated in such a non-exciting experiment as Experiment 1, they can be considered practically identifiable.

Experiment 2.

The performance of the Monte Carlo analysis for the 4 different noise levels, i.e., 5%, 10%, 15% and 25% is reported in Table 4. The successful parameter estimations took about 25-45 minutes to solve. The time needed in unsuccessful parameter estimation varied between about 5-45 minutes.

Table 4: Performance of the Monte Carlo analysis for experiment 2.

Noise	R_1	R_2	R_3
5%	4.2813	4.2507	0.9939
10%	6.3492	6.1943	0.9920
15%	5.1050	4.8934	0.9932
25%	4.8305	5.0734	0.9923

Because $R_3 \approx 1$, the data on which the Monte Carlo analysis was performed, is assumed to be compliant with the assumptions concerning random sampling. Generally speaking the number of failed parameter estimations is higher for Experiment 2 than for Experiment 1. Comparing R_1 and R_2 in Table 4 with R_1 and R_2 in Table 3 clearly shows that the quality of the estimates is much lower than for Experiment 1. The error is several factors larger than the measurement errors. A possible explanation for this error is a bias in the parameter estimations. To check if there is a bias in the parameter estimations, the average normalized mean error between the predictions and the samples

Noise level	S_1	S_2
5%	0.6028	0.6033
10%	0.5542	0.5542
15%	0.6002	0.5990
25%	0.4384	0.4374

Table 5: S_1 and S_2 of the Monte Carlo analysis for experiment 2.

(S_1) and the true values (S_2), introduced in Equation (31) and (32), were calculated. For an accurate parameter estimation, both S_1 and S_2 should be around zero. The results for the different noise levels are listed in Table 5

$$S_1 = \frac{1}{n_{it}} \frac{1}{n_t} \frac{1}{n_y} \sum_{i=1}^{n_{it}} \sum_{i_t=1}^{n_t} \sum_{i_y=1}^{n_y} \frac{(y_{i_y, i_t} - y_{i_y, i_t}^s)}{\sigma_{i_y}} \quad (31)$$

$$S_2 = \frac{1}{n_{it}} \frac{1}{n_t} \frac{1}{n_y} \sum_{i=1}^{n_{it}} \sum_{i_t=1}^{n_t} \sum_{i_y=1}^{n_y} \frac{(y_{i_y, i_t} - y_{i_y, i_t}^*)}{\sigma_{i_y}} \quad (32)$$

It is clear that there is a positive bias in the results: the prediction values are generally higher than the true values or measurements. This is due to the fact that the values of some states almost become zero. See for instance the profile of S_{nh3} in Figure 8. During the acidification of the system, the value drops from $4.5 \cdot 10^{-3}$ to about 10^{-6} . Since the noise is normally distributed with a standard deviation equal to about 5%-25% of $4.5 \cdot 10^{-3}$, negative measurement values are ‘observed’. Hence, there could be Monte Carlo realizations with measurements such that the solution that maximizes the likelihood as defined in Equation (17), goes below zero. These solutions, however, are forbidden due to the imposed lower bounds. Therefore solutions will be selected that have only positive states and are, hence, positively biased in comparison to the measurements and the true values. It is believed that this parameter estimation behavior is also responsible for the large number of failed parameter estimates.

656 This problem cannot be solved in straightforward way. Either the assump-
 657 tions on the noise are changed by making them for instance proportional to
 658 the value of the state, or the biased estimates are used. The latter option is
 659 chosen since the former implies a change of the practical experimental condi-
 660 tions.

661 The parameter estimates results for the successful experiments are listed in
 662 Table 3 and 4 of Appendix E in which the bias and the observed standard de-
 663 viation are reported. Note that because of the pH-control with cations $S_{\text{cat}^+, \text{in}}$
 664 and anions $S_{\text{an}^-, \text{in}}$, the states S_{cat^+} and S_{an^-} are no longer included in the
 665 model. Thus, the initial conditions $S_{\text{cat}^+}(0)$ and $S_{\text{an}^-}(0)$ are no longer esti-
 666 mated. The total number of parameters and unknowns for this Experiment is
 667 thus 98. The results of Table 3 and 4 are discussed in detail in Appendix E.

668 In terms of relative parameter bias (cfr. Equation (21)) and parameter stan-
 669 dard deviation (cfr. Equation (22)), (and considering the successful experi-
 670 ments) the results are generally better than for Experiment 1. For instance,
 671 for the case of 5% noise, there are 92 ‘good’ entries, 69 ‘moderate’ entries and
 672 39 ‘bad’ entries, i.e., the relative bias or standard deviation is lower than 5%,
 673 between 5% and 25% and higher than 25%, respectively. This is better than
 674 the 86 ‘good’ entries, 42 ‘moderate’ and 72 ‘bad’ entries for Experiment 1.

675 It is apparent that the parameters associated with the states, i.e., the stoichio-
 676 metric, composition, kinetic, and decay parameters, that were directly excited,
 677 i.e., S_{su} and S_{aa} are estimated well. This is in addition to the parameters that
 678 were already found to be practically identifiable in Experiment 1. The other
 679 parameters associated with the other states are also better estimated, with
 680 the exception for the parameters associated with the degradation of S_{fa} and

681 S_{ac} . Especially for the latter, the estimates are very poor. Thus, the idea be-
 682 hind the heuristic design, i.e., a sufficiently large domain of state values, is
 683 not sufficient to guarantee practical identifiability for the all non-linear ex-
 684 pressions. A clue to a better experiment design is the observation that the di-
 685 rect excitation of the states through the change of input, resulted in ‘sharper’
 686 peaks in the concentration profile of S_{su} and S_{aa} than for the other states (see
 687 Figure 8).

688 To investigate the interdependence between the estimates, the normalized co-
 689 variance matrix as defined in Equation (23) is calculated and depicted in Fig-
 690 ure 9. Only the result for the 5% noise level is analyzed. Again, values below
 691 5 % are considered to correspond with weak or no relation between the pa-
 692 rameter, values between 5% and 25% correspond to a moderate relation and
 693 values higher than 25% indicate a strong relation between the parameters.

694 There are 2720 pairs with no or weak interdependence, 1901 pairs with mod-
 695 erate interdependence and 294 pairs of strongly related parameter estimates in
 696 the upper triangle of the matrix. This does not include the diagonal elements
 697 of the matrix which represent absolute values of the single parameter relative
 698 standard deviations. In comparison with the first experiment, there are less
 699 strong interdependencies. However, there a bit more moderate interdependen-
 700 cies.

701 Generally, the results of the covariance matrix correspond with what was con-
 702 cluded on the results of Tables 3 and 4 of Appendix E. In this experiment, in
 703 addition to the parameters that were already found to be practically identifi-
 704 able in the first experiment, the parameters concerning the S_{aa} and S_{su} show
 705 little interdependency with other parameters.

706 In conclusion, this informative experiment showed that through the direct ex-
707 citation of the mono-saccharides and amino acids, the associated degradation
708 kinetic, stoichiometric and composition parameters are practically identifiable
709 in the stated experimental conditions.

710 If the results of the two experiments are combined, it shows that at least the
711 first-order kinetic parameters (36.-40.), the stoichiometric parameters associ-
712 ated with the degradation of lipids (1.), monosaccharides (14.-17.) and amino
713 acids (18.-22.), the composition of amino acids and monosaccharides (29., 31.,
714 33), two of the 3 pH-inhibition level pairs (66.-67., 70.-71.) and the saturation
715 constant for acetic acid degradation (55.) are practically identifiable. This
716 does not mean that the other parameters are *not* practically identifiable, just
717 that they were not practically identifiable in the designed experiment. Based
718 on the results described in this manuscript, there is still room for improve-
719 ment in the design of the experiment. For instance, all states can directly be
720 excited, rather than exciting a selected few and relying on the interconnectiv-
721 ity of the model to excite the other states indirectly.

722 Also, the results are specific for the considered set of true parameter values,
723 the nature of the noise, the choice of available inputs and outputs and under
724 the assumption that the anaerobic digestion system is in reality described by
725 the ADM1. Nevertheless, they are indicative on what can be expected from a
726 parameter estimation procedure on real data.

727 5. Conclusions

728 In this work, the structural and practical identifiability of the Anaerobic Di-
729 gestion Model No. 1 is analyzed as an example of a large-scale non-linear bi-
730 ological network model, consisting of algebraic equations and non-rational ex-
731 pressions.

732 The authors have presented modifications to the model, in terms of parameter
733 combinations of structurally nonidentifiable parameters to yield an as good as
734 fully locally structurally identifiable model structure. The *probabilistic* algo-
735 rithm of [20] has been applied to investigate the structural identifiability for
736 different input and output scenarios.

737 Furthermore, the generally positive structural identifiability results can be ex-
738 plained from the large number of interconnections between the states in the
739 network structure. This interconnectivity, however, is also observed in the
740 parameter estimates, making uncorrelated parameter estimations in practice
741 difficult. Structural non-identifiability occurs at structural dead-ends in the
742 ADM1 network as the inerts and for the anions/cations which are only ap-
743 pearing in pairs in the equations.

744 The practical identifiability of the model is analyzed by solving repeatedly
745 large-scale parameter estimation problems. Based on a non-informative and
746 an informative experiment the practical identifiability of the ADM1 case study
747 was analyzed. At least the following parameters were found to be practically
748 identifiable: (i) the first-order kinetic parameters except the biomass decay
749 constants, (ii) the stoichiometric parameters associated with the degradation
750 of lipids, monosaccharides, and amino acids, (iii) the composition of amino

751 acids and monosaccharides, (*iv*) two of the 3 pH-inhibition levels and (*v*) the
 752 saturation constant for acetic acid degradation.

753 Generally speaking it is found that it is better to directly excite the states by
 754 means of the input, than relying on the interactions of the biological network
 755 to indirectly excite the states of interest. Additionally, the case study shows
 756 a strong interconnectivity between the states. This interconnectivity is also
 757 observed in the covariances of the parameter estimates. Therefore, it is not
 758 recommended to estimate parameters as if the reactions are completely sepa-
 759 rated.

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767 References

- 768 [1] D. Batstone, J. Keller, I. Angelidaki, S. Kalyuzhnyi, S. Pavlostathis,
 769 A. Rozzi, W. Sanders, H. Siegrist, V. Vavilin, The IWA anaerobic diges-
 770 tion model No 1 (ADM1), Water Science and Technology 45 (10) (2002)
 771 65–73.

- 772 [2] D. J. Batstone, J. Keller, I. Angelidaki, Anaerobic Digestion Model No.
773 1, IWA Publishing, 2002.
- 774 [3] F. Blumensaat, J. Keller, Modelling of two-stage anaerobic digestion us-
775 ing the IWA anaerobic digestion model no. 1 (ADM1), Water Research
776 39 (1) (2005) 171–183.
- 777 [4] S. Weinrich, M. Nelles, Critical comparison of different model structures
778 for the applied simulation of the anaerobic digestion of agricultural en-
779 ergy crops, Bioresource Technology 178 (2015) 306 – 312.
- 780 [5] N. Kythreotou, G. Florides, S. A. Tassou, A review of simple to scientific
781 models for anaerobic digestion, Renewable Energy 71 (2014) 701 – 714.
- 782 [6] C. Mendes, K. Esquerre, L. M. Queiroz, Application of anaerobic diges-
783 tion model no. 1 for simulating anaerobic mesophilic sludge digestion,
784 Waste Management 35 (2015) 89 – 95.
- 785 [7] C. Rosen, U. Jeppsson, Aspects on ADM1 implementa-
786 tion within the BSM2 framework, [Online]. Available:
787 <http://ftp.iea.lth.se/publications/Reports/LTH-IEA-7224.pdf> (2006).
- 788 [8] K. Koch, M. Lübken, T. Gehring, M. Wichern, H. Horn, Biogas from
789 grass silage–measurements and modeling with ADM1, Bioresource Tech-
790 nology 101 (21) (2010) 8158–8165.
- 791 [9] F. Mairet, O. Bernard, M. Ras, L. Lardon, J.-P. Steyer, Modeling anaero-
792 bic digestion of microalgae using ADM1, Bioresource Technology 102 (13)
793 (2011) 6823–6829.
- 794 [10] I. Ntaikou, H. N. Gavala, G. Lyberatos, Application of a modified anaer-
795 obic digestion model 1 version for fermentative hydrogen production from

- 796 sweet sorghum extract by ruminococcus albus, International Journal of
 797 Hydrogen Energy 35 (8) (2010) 3423 – 3432.
- 798 [11] V. Razaviarani, I. D. Buchanan, Calibration of the anaerobic digestion
 799 model no. 1 (adm1) for steady-state anaerobic co-digestion of municipal
 800 wastewater sludge with restaurant grease trap waste, Chemical Engineer-
 801 ing Journal 266 (2015) 91 – 99.
- 802 [12] R. Kleerebezem, M. Van Loosdrecht, Waste characterization for imple-
 803 mentation in ADM1, Biotechnology and Bioengineering 54 (4) (2006)
 804 167–174.
- 805 [13] U. Zaher, P. Buffiere, J.-P. Steyer, S. Chen, A procedure to estimate
 806 proximate analysis of mixed organic wastes, Water Environment Research
 807 81 (2009) 407–415.
- 808 [14] H. Junicke, B. Abbas, J. Oentoro, M. van Loosdrecht, M. Kleere-
 809 bezem, Absolute quantification of individual biomass concentrations in
 810 a methanogenic coculture, AMB express 35 (4) (2014) 1–8.
- 811 [15] E. Walter, L. Pronzato, Identification of Parametric Models from Experi-
 812 mental Data, Springer Verlag, Heidelberg, 1997.
- 813 [16] K. Godfrey, J. DiStefano, Identifiability of model parameters, in: E. Wal-
 814 ter (Ed.), Identifiability of Parametric Models, Pergamon Press, Oxford,
 815 1987, pp. 1–20.
- 816 [17] H. Pohjanpalo, System identifiability based on the power series expansion
 817 of the output, Mathematical Biosciences 41 (1) (1978) 21–33.
- 818 [18] M. Saccomani, S. Audoly, L. D’Angio, Parameter identifiability of non-
 819 linear systems: the role of initial conditions, Automatica 39 (4) (2003)
 820 619–632.

- 821 [19] O. Chiş, J. R. Banga, E. Balsa-Canto, GenSSI: a software toolbox for
 822 structural identifiability analysis of biological models, *Bioinformatics*
 823 27 (18) (2011) 2610–2611.
- 824 [20] A. Sedoglavic, A probabilistic algorithm to test local algebraic observability
 825 in polynomial time, *Journal of Symbolic Computation* 33 (2002) 735–755.
- 826 [21] J. Karlsson, M. Anguelova, M. Jirstrand, An efficient method for struc-
 827 tural identifiability analysis of large dynamic systems, in: *Procs of the*
 828 *16th IFAC Symposium on System Identification, Brussels (Belgium),*
 829 *2012, pp. 941–946.*
- 830 [22] M. Anguelova, Observability and identifiability of nonlinear systems
 831 with applications in biology, Ph.D. thesis, Dept. Mathematical Science,
 832 Chalmers University of Technology and Göteborg University, Göteborg
 833 (2007).
- 834 [23] S. Diop, Y. Wang, Equivalence between algebraic observability and local
 835 generic observability, in: *Proceedings of the 32nd Conference on Decision*
 836 *and Control, Vol. 3, San Antonio, TX (USA), 1993, pp. 2864 – 2865.*
- 837 [24] R. Hermann, A. J. Krener, Nonlinear controllability and observability,
 838 *IEEE Transactions on Automatic Control* 22 (5) (1977) 728–740.
- 839 [25] E. Sontag, A concept of local observability, *Systems and Control Letters*
 840 5 (1) (1984) 41–47.
- 841 [26] A. Raue, J. Karlsson, M. Saccomani, M. Jirstrand, T. J., Comparison of
 842 approaches for parameter identifiability analysis of biological systems,
 843 *Bioinformatics* 30 (10) (2014) 1140–1148.
- 844 [27] G. Bellu, M. P. Saccomani, S. Audoly, L. D’Angiò, DAISY: a new soft-
 845 ware tool to test global identifiability of biological and physiological sys-

tems, Computer methods and programs in biomedicine 88 (1) (2007) 52–
61.

[28] C. D. Knightes, C. A. Peters, Statistical analysis of nonlinear parameter estimation for monod biodegradation kinetics using bivariate data, Biotechnology and bioengineering 69 (2) (2000) 160–170.

[29] T. G. Ellis, D. S. Barbeau, B. F. Smets, C. L. Grady Jr, Respirometric technique for determination of extant kinetic parameters describing biodegradation, Water Environment Research 68 (5) (1996) 917–926.

[30] A. Holmberg, On the practical identifiability of microbial growth models incorporating michaelis-menten type nonlinearities, Mathematical Biosciences 62 (1) (1982) 23–43.

[31] S. S. Bhonsale, D. Telen, D. Vercammen, M. Vallerio, J. Hufkens, P. Nimmegeers, F. Logist, J. Van Impe, Pomodoro - an open source toolkit for multiobject optimal control, and model based control and estimation, Preprint submitted to Expert Systems with Applications.

[32] S. Bhonsale, M. Vallerio, D. Telen, D. Vercammen, F. Logist, J. Van Impe, SolACE: An open source package for nonlinear model predictive control and state estimation for (bio)chemical processes, in: Proceedings of the 26th European Symposium on Computer Aided Process Engineering. Portoroz, Slovenia., June 12th - 15th, 2016.

[33] L. T. Biegler, Nonlinear Programming: Concepts, Algorithms, and Applications to Chemical Processes, Society for Industrial and Applied Mathematics (Philadelphia, USA), 2010, Chapter 10: Simultaneous Methods for Dynamic Optimization.

[34] L. A. Alves, J. B. Almeida e Silva, M. Giuliatti, Solubility of d-glucose in

- 871 water and ethanol/water mixtures, Journal of Chemical & Engineering
872 Data 52 (6) (2007) 2166–2170.
- 873 [35] D. Strik, A. Domnanovich, P. Holubar, A pH-based control of ammonia
874 in biogas during anaerobic digestion of artificial pig manure and maize
875 silage, Process Biochemistry 41 (6) (2006) 1235–1238.

876 Figures

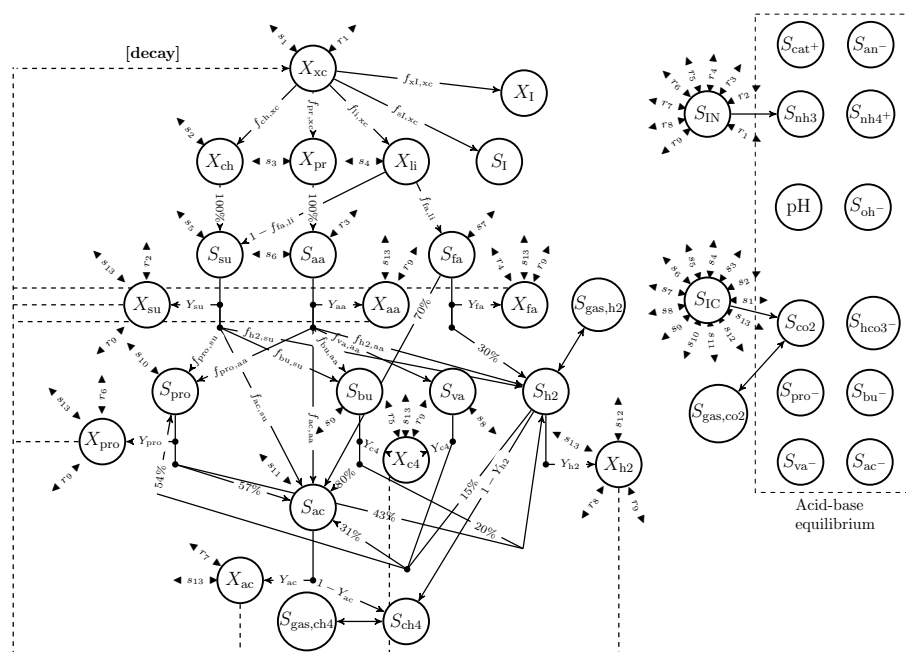


Figure 1: **Reaction network of ADM1.** The structure of ADM1 [1], on which the stoichiometric parameters $f_{i,j}$ and Y_i are indicated. Inorganic nitrogen S_{IN} and carbon S_{IC} are source/sink components that collect or make up excess or deficiencies of carbon and nitrogen, due to different N- or C-compositions of the different components in the system, indicated with r_i and s_i respectively. A separate system defines the acid-base equilibrium. The microbial metabolism on the substrates is inhibited by the pH-level, limiting inorganic nitrogen, ammonia and hydrogen.

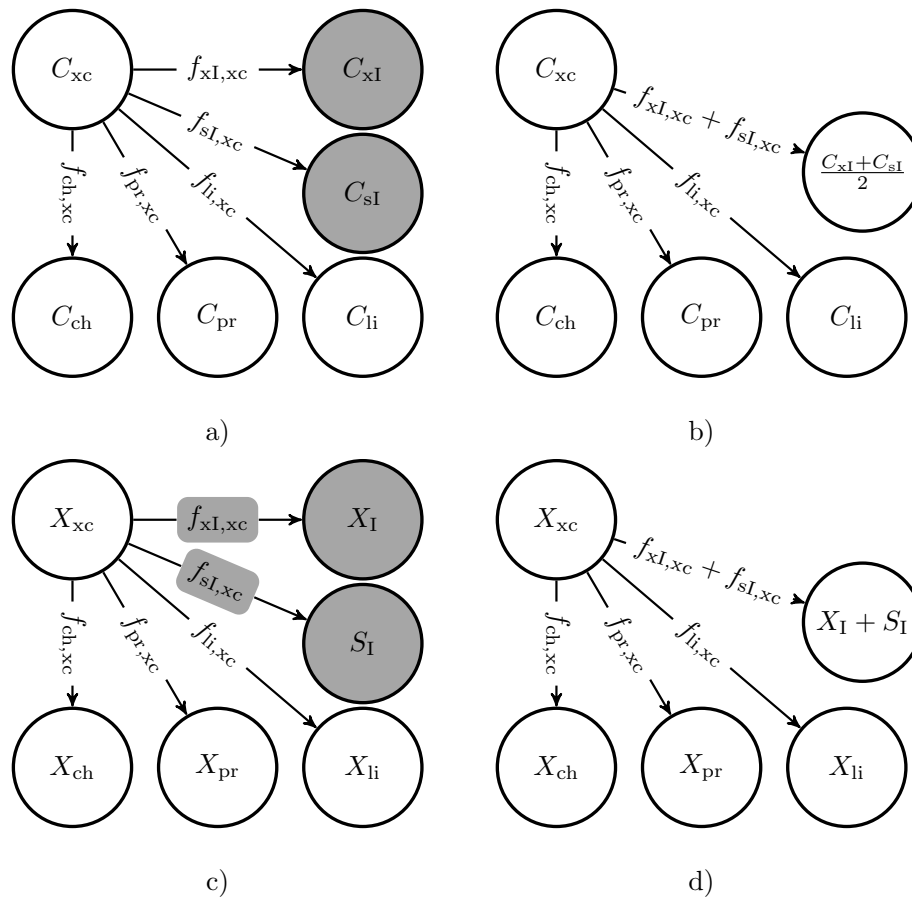


Figure 2: **Proposed parameter combinations for ADM1.** The distribution of carbon from the degradation of X_{xc} as described in a) the original equations of ADM1 and b) with the proposed parameter combination. The distribution of COD when none of the inerts are measured as described in c) the original equations of ADM1 and d) with the proposed parameter combination. Dark gray indicates unidentifiability.

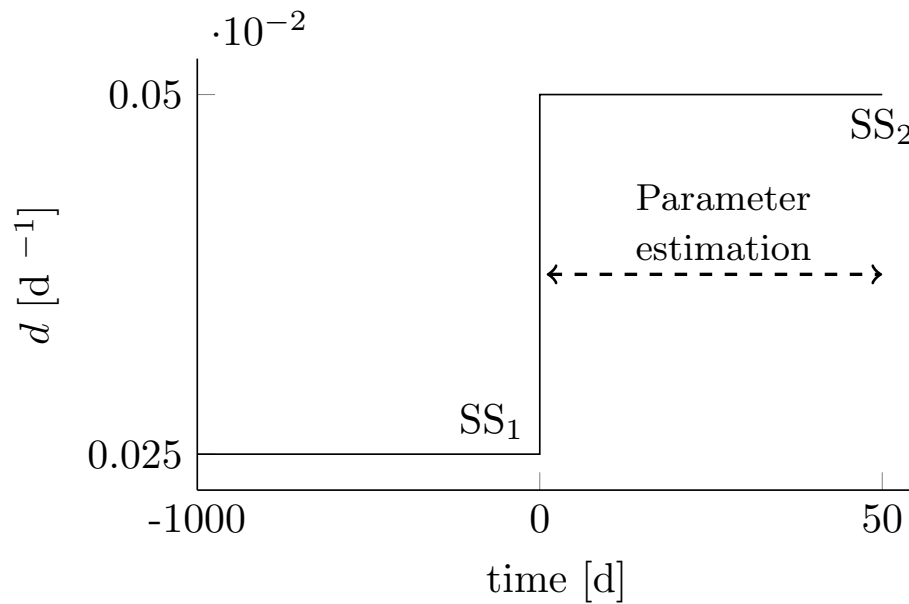


Figure 3: **Dilution rate d as a function of time for Experiment 1.** The profile starts from a steady-state (SS_1) obtained by applying a dilution rate of 0.025 days^{-1} for 1000 days. The dilution rate is then increased to 0.05 days^{-1} for 50 days. Eventually the system will reach a new steady-state SS_2 . The profile considered for the parameter estimation is indicated by the double arrow.

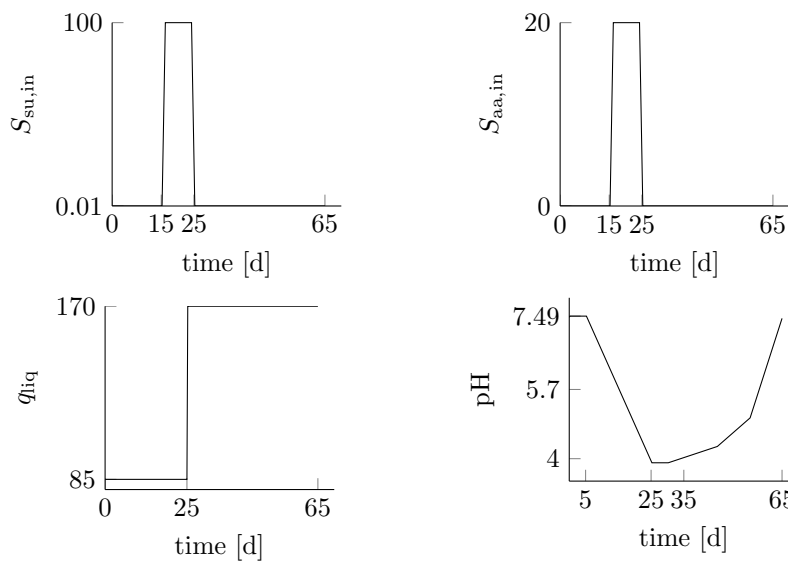


Figure 4: **Applied input for Experiment 2.** Input concentrations of $S_{su,in}$, $S_{an,in}$, q_{liq} and the desired pH profile of experiment 2.

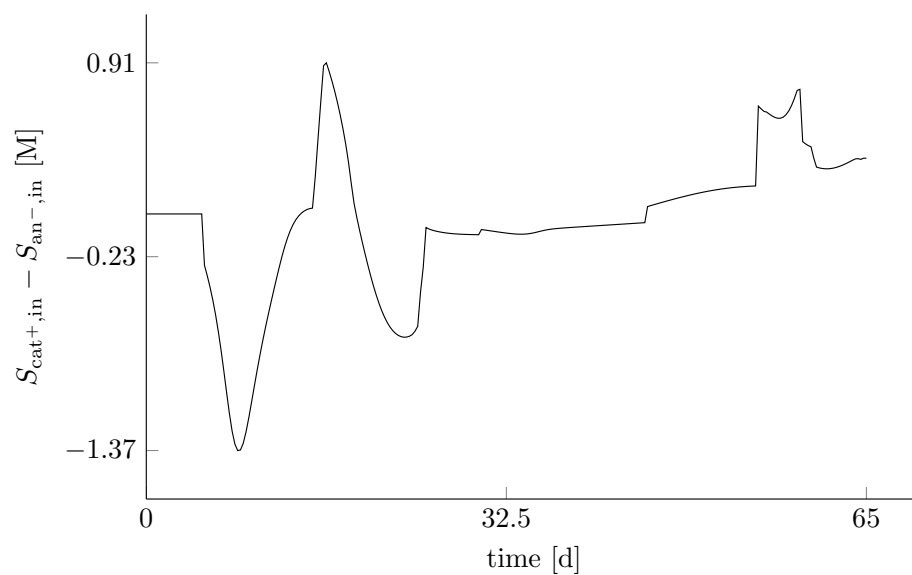


Figure 5: **The applied** $S_{\text{cat}^+, \text{in}} - S_{\text{an}^-, \text{in}}$ profile needed to obtain the pH-profile of Figure 4.

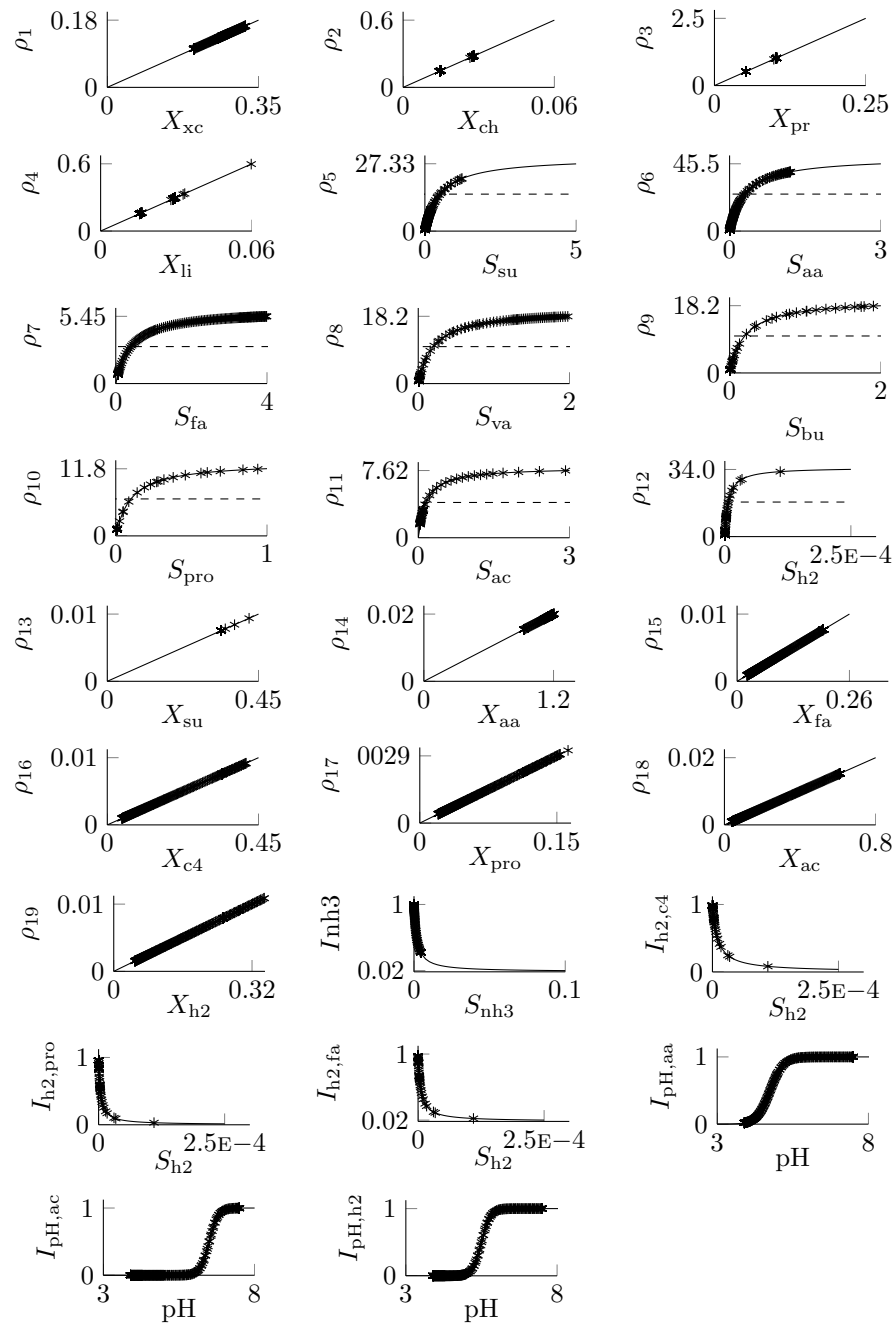


Figure 6: Plot of the simulation results of Experiment 2 on the reaction rates of ADM1. The value $\mu_{max}/2$ is indicated with a dashed line (- -).

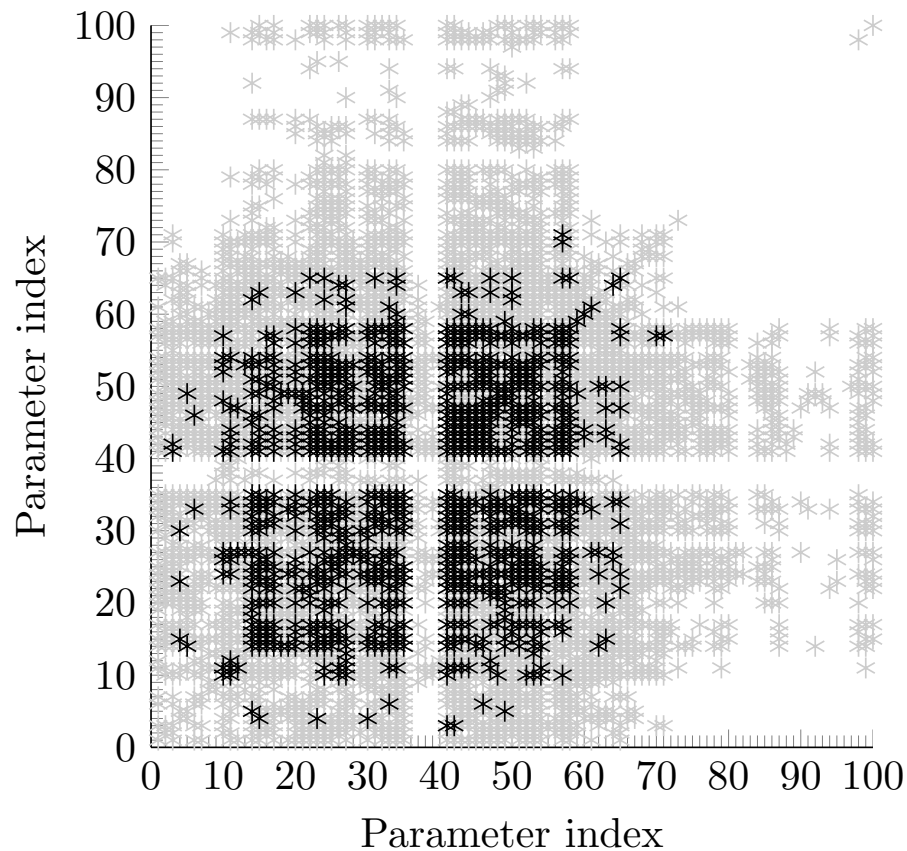


Figure 7: **Square root of the absolute relative covariances for the parameters estimates at a 5% noise level for Experiment 1.** The parameter numbering of Appendix E is used. A root of an absolute covariance higher than 25% is indicated in black. A root of an absolute covariance in between 5% and 25% is indicated in grey. Roots of absolute covariances lower than 5 % are not indicated.

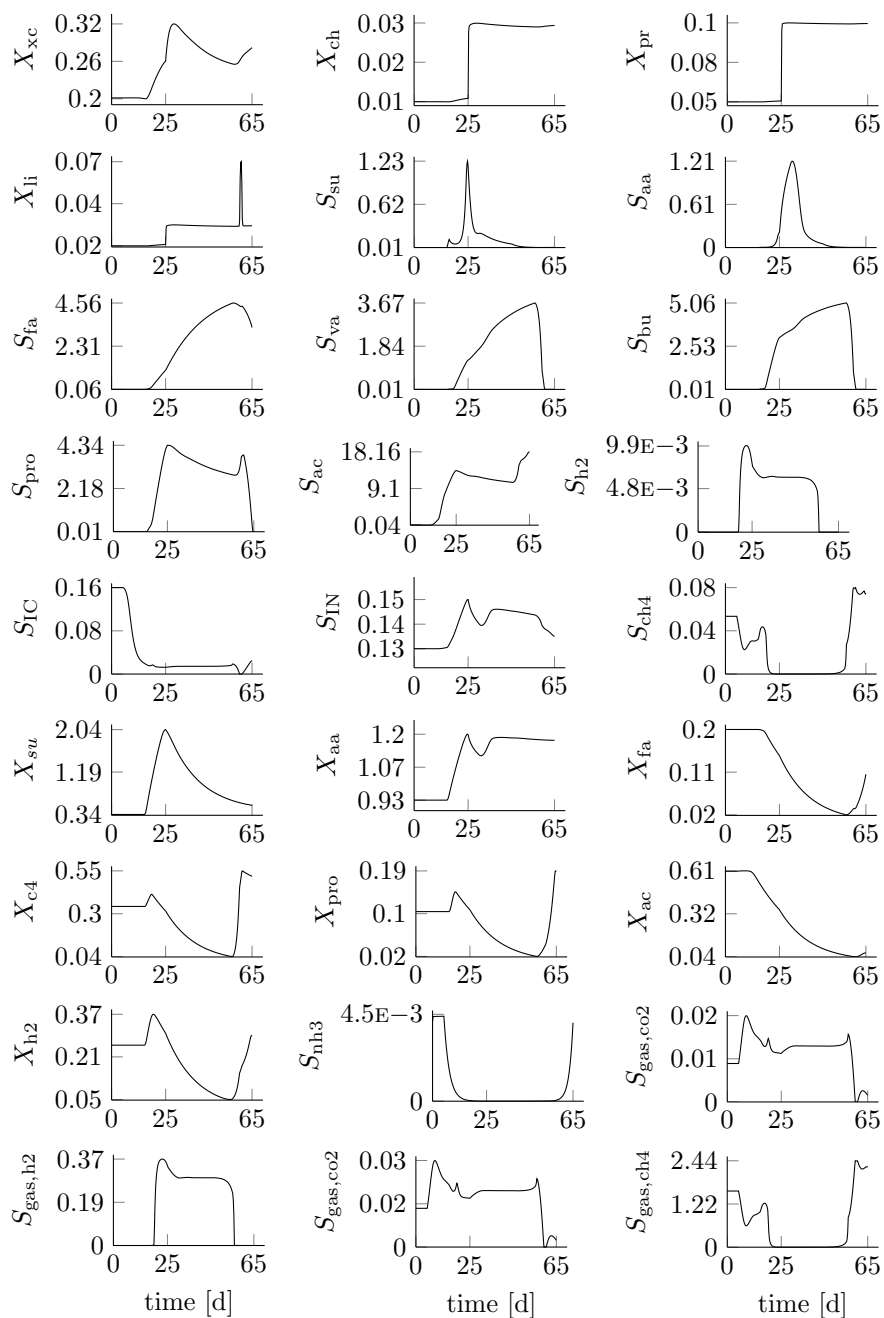


Figure 8: Concentration profiles of most states in Experiment 2.

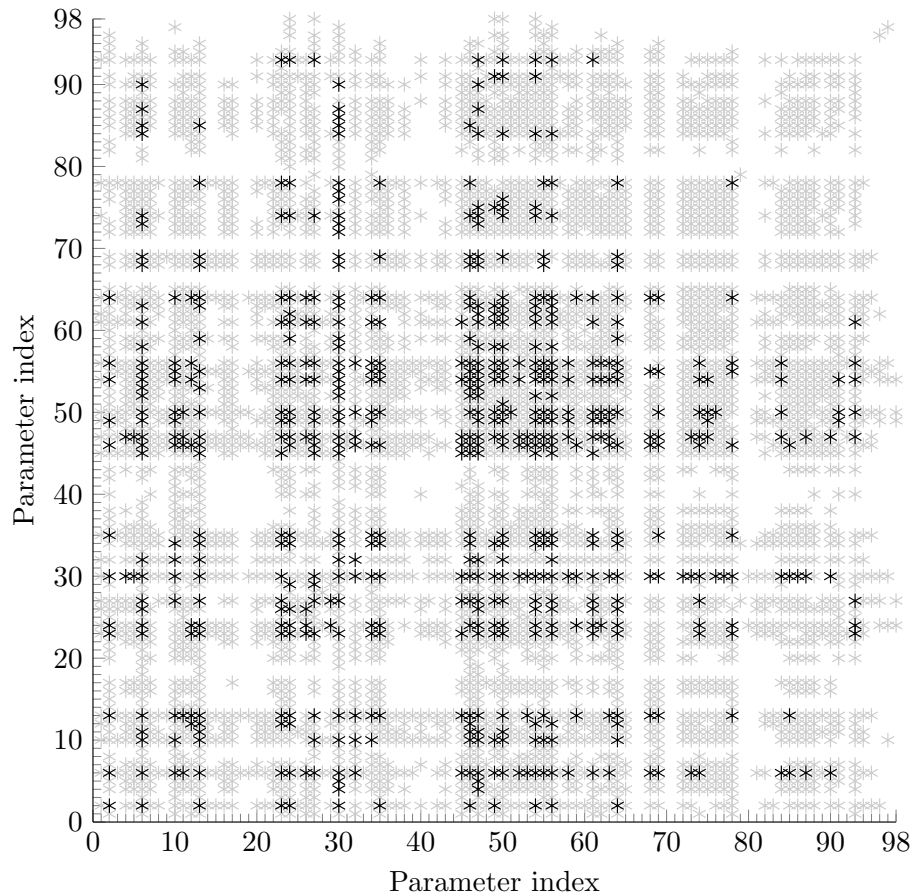


Figure 9: **Square root of the absolute relative covariances for the parameters estimates at a 5% noise level for Experiment 2.** The parameter numbering of Appendix E is used. A root of an absolute covariance higher than 25% is indicated in black. A root of an absolute covariance in between 5% and 25% is indicated in gray. Roots of absolute covariances lower than 5 % are not indicated.

877 **Appendix A. Description of ADM1**

878 **Appendix B. Numerical values of states and parameters values**

879 **Appendix C. Relations between ADM1 model structure and struc-**
880 **tural identifiability results**

881 **Appendix D. In-depth discussion of the state profiles in Experi-**
882 **ment 2**

883 **Appendix E. Parameter estimation results**

Appendix A. Description of ADM1

In this appendix, the states, parameters, reactions and equations of ADM1 that are used in this article are listed. The state vector and parameter vector given by \mathbf{x} and $\boldsymbol{\theta}$.

$$\mathbf{x} = \left\{ \begin{array}{c} S_{su} \\ S_{aa} \\ S_{fa} \\ S_{va} \\ S_{bu} \\ S_{pro} \\ S_{ac} \\ S_{h2} \\ S_{ch4} \\ S_{IC} \\ S_{IN} \\ S_I \\ X_{xc} \\ X_{ch} \\ X_{pr} \\ X_{li} \\ X_I \\ X_{su} \\ X_{aa} \\ X_{fa} \\ X_{c4} \\ X_{pro} \\ X_{ac} \\ X_{h2} \\ S_{gas,h2} \\ S_{gas,ch4} \\ S_{gas,co2} \\ S_{cat+} \\ S_{an-} \\ S_{va-} \\ S_{bu-} \\ S_{pro-} \\ S_{ac-} \\ S_{oh-} \\ S_{h+} \\ S_{hco3-} \\ S_{co2} \\ S_{nh4+} \\ S_{nh3} \end{array} \right\} \quad (1)$$

States

Table 1: Overview of the states.

State	Description	Unit
S_{su}	Soluble monosaccharides	kg COD m ⁻³
S_{aa}	Soluble amino acids	kg COD m ⁻³
S_{fa}	Soluble long chain fatty acids	kg COD m ⁻³
S_{va}	Soluble valeric acid	kg COD m ⁻³
S_{bu}	Soluble butyric acid	kg COD m ⁻³
S_{pro}	Soluble propionic acid	kg COD m ⁻³
S_{ac}	Soluble acetic acid	kg COD m ⁻³
S_{h2}	Soluble hydrogen	kg COD m ⁻³
S_{ch4}	Soluble methane	kg COD m ⁻³
S_{IC}	Soluble inorganic carbon	kmol C m ⁻³
S_{IN}	Soluble inorganic nitrogen	kmol N m ⁻³
S_I	Soluble inerts	kg COD m ⁻³
X_{xc}	Particulate composites	kg COD m ⁻³
X_{ch}	Particulate carbohydrates	kg COD m ⁻³
X_{pr}	Particulate proteins	kg COD m ⁻³
X_{li}	Particulate lipids	kg COD m ⁻³
X_I	Particulate inerts	kg COD m ⁻³
X_{su}	Monosaccharide degraders (biomass)	kg COD m ⁻³
X_{aa}	Amino acid degraders (biomass)	kg COD m ⁻³
X_{fa}	Fatty acid degraders (biomass)	kg COD m ⁻³
X_{c4}	Valeric and butyric acid degraders (biomass)	kg COD m ⁻³
X_{pro}	Propionic acid degraders (biomass)	kg COD m ⁻³
X_{ac}	Acetic acid degraders (biomass)	kg COD m ⁻³
X_{h2}	Hydrogen degraders (biomass)	kg COD m ⁻³
$S_{gas,h2}$	Gaseous hydrogen	kg COD m ⁻³
$S_{gas,ch4}$	Gaseous methane	kg COD m ⁻³
$S_{gas,co2}$	Gaseous carbon dioxide	kg COD m ⁻³
S_{cat+}	Cations	kmol m ⁻³
S_{an-}	Anions	kmol m ⁻³
S_{va-}	Valerate	kg COD m ⁻³
S_{bu-}	Butyrate	kg COD m ⁻³
S_{pro-}	Propionate	kg COD m ⁻³
S_{ac-}	Acetate	kg COD m ⁻³
S_{oh-}	Hydroxide	kmol H ⁺ m ⁻³
S_{h+}	Proton	kg COD m ⁻³
S_{hco3-}	Bicarbonate	kmol C m ⁻³
S_{co2}	Soluble carbon dioxide	kmol C m ⁻³
S_{nh4+}	Ammonium	kmol N m ⁻³
S_{nh3}	Ammonia	kmol N m ⁻³

Reactions

Table 2: Reaction rates.

Symbol	Description	Symbol	Description
ρ_1	Disintegration of X_{xc}	ρ_{12}	Methanogenesis of S_{h2}
ρ_2	Hydrolysis of X_{ch}	ρ_{13}	Decay of X_{su}
ρ_3	Hydrolysis of X_{pr}	ρ_{14}	Decay of X_{aa}
ρ_4	Hydrolysis of X_{li}	ρ_{15}	Decay of X_{fa}
ρ_5	Acidogenesis of S_{su}	ρ_{16}	Decay of X_{c4}
ρ_6	Acidogenesis of S_{aa}	ρ_{17}	Decay of X_{pro}
ρ_7	Acidogenesis of S_{fa}	ρ_{18}	Decay of X_{ac}
ρ_8	Acetogenesis of S_{va}	ρ_{19}	Decay of X_{h2}
ρ_9	Acetogenesis of S_{bu}	$\rho_{T,8}$	liquid-gas transfer S_{h2}
ρ_{10}	Acetogenesis of S_{pro}	$\rho_{T,9}$	liquid-gas transfer S_{ch4}
ρ_{11}	Methanogenesis S_{ac}	$\rho_{T,10}$	liquid-gas transfer S_{co2}

Parameters

Table 3: Stoichiometric parameters.

Symbol	Description	Symbol	Description
$f_{sl,xc}$	Yield of S_I on X_{xc}	$f_{pro,aa}$	Yield of S_{pro} on S_{aa}
$f_{ch,xc}$	Yield of X_{ch} on X_{xc}	$f_{va,aa}$	Yield of S_{va} on S_{aa}
$f_{pr,xc}$	Yield of X_{pr} on X_{xc}	$f_{ac,su}$	Yield of S_{ac} on S_{su}
$f_{xl,xc}$	Yield of X_I on X_{xc}	$f_{pro,su}$	Yield of S_{pro} on S_{su}
$f_{li,xc}$	Yield of X_{li} on X_{xc}	$f_{h2,su}$	Yield of S_{h2} on S_{su}
$f_{ac,aa}$	Yield of S_{ac} on S_{aa}	$f_{bu,su}$	Yield of S_{bu} on S_{su}
$f_{bu,aa}$	Yield of S_{bu} on S_{aa}	$f_{fa,li}$	Yield of S_{fa} on X_{li}
$f_{h2,aa}$	Yield of S_{h2} on S_{aa}		
Y_{su}	Yield of X_{su}	Y_{pro}	Yield of X_{pro}
Y_{aa}	Yield of X_{aa}	Y_{c4}	Yield of X_{c4}
Y_{fa}	Yield of X_{fa}	Y_{h2}	Yield of X_{h2}
Y_{ac}	Yield of X_{ac}		

Table 4: Composition parameters.

Symbol	Description	Symbol	Description
C_{aa}	C content S_{aa}	C_{pr}	C content X_{pr}
C_{ac}	C content S_{ac}	C_{pro}	C content S_{pro}
C_{bac}	C content X_{su}	C_{sl}	C content S_I
C_{bu}	C content S_{bu}	C_{su}	C content X_{ch}
C_{ch}	C content X_{ch}	C_{va}	C content S_{va}

C_{ch4}	C content S_{ch4}	C_{xc}	C content X_{xc}
C_{fa}	C content S_{fa}	C_{xI}	C content X_I
C_{li}	C content X_{li}		
N_{aa}	N content S_{aa} and X_{pr}	N_I	N content S_I and X_I
N_{bac}	N content X_{su} , X_{aa} , X_{fa} , X_{ac} , X_{pro} , X_{c4} and X_{h2}	N_{xc}	N content X_{xc}

Table 5: Kinetic parameters.

Symbol	Description	Symbol	Description
k_{dis}	First-order ρ_1	$k_{dec,Xaa}$	First-order ρ_{14}
$k_{hyd,ch}$	First-order ρ_2	$k_{dec,Xfa}$	First-order ρ_{15}
$k_{hyd,pr}$	First-order ρ_3	$k_{dec,Xc4}$	First-order ρ_{16}
$k_{hyd,li}$	First-order ρ_4	$k_{dec,Xpro}$	First-order ρ_{17}
$k_{m,su}$	Maximum rate ρ_5	$k_{dec,Xac}$	First-order ρ_{18}
$K_{S,su}$	Saturation ρ_5	$k_{dec,Xh2}$	first-order ρ_{19}
$k_{m,aa}$	Maximum rate ρ_6	$K_{IN,lim}$	Limiting-N inhib. $\rho_5 - \rho_{12}$
$K_{S,aa}$	Saturation ρ_6	$K_{I,h2,fa}$	H ₂ -inhibition ρ_7
$k_{m,fa}$	Maximum rate ρ_8	$K_{I,h2,c4}$	H ₂ -inhibition ρ_8 and ρ_9
$K_{S,fa}$	Saturation ρ_8	$K_{I,h2,pro}$	H ₂ -inhibition ρ_{10}
$k_{m,c4}$	Maximum rate ρ_9 and ρ_{10}	$K_{I,nh3}$	NH ₃ -inhibition ρ_{11}
$K_{S,c4}$	Saturation ρ_9 and ρ_{10}	$pH_{LL,aa}$	Lower limit pH-inhib. $\rho_5 - \rho_{10}$
$k_{m,pro}$	Maximum rate ρ_{10}	$pH_{UL,aa}$	Upper limit pH-inhib. $\rho_5 - \rho_{10}$
$K_{S,pro}$	Saturation ρ_{10}	$pH_{LL,h2}$	Lower limit pH-inhib. ρ_{11}
$k_{m,ac}$	Maximum rate ρ_{11}	$pH_{UL,h2}$	Upper limit pH-inhib. ρ_{11}
$K_{S,ac}$	Saturation ρ_{11}	$pH_{LL,ac}$	Lower limit pH-inhib. ρ_{12}
$k_{m,h2}$	Maximum rate ρ_{12}	$pH_{UL,ac}$	Upper limit pH-inhib. ρ_{12}
$K_{S,h2}$	Saturation ρ_{12}	k_{La}	Liquid-gas mass-transfer
$k_{dec,Xsu}$	First-order ρ_{13}		

Table 6: Physico-chemical constants.

Symbol	Description	Symbol	Description
K_w	Dissociation constant water	$K_{H,co2}$	Henry constant S_{co2}
$K_{a,va}$	Dissociation constant S_{va}	$K_{H,h2}$	Henry constant S_{h2}
$K_{a,bu}$	Dissociation constant S_{bu}	$K_{H,ch4}$	Henry constant S_{ch4}
$K_{a,pro}$	Dissociation constant S_{pro}	R	Universal gas constant
$K_{a,ac}$	Dissociation constant S_{ac}	p_{h2o}	Vapor pressure water
$K_{a,co2}$	Dissociation constant S_{co2}	p_{atm}	Atmospheric pressure
$K_{a,IN}$	Dissociation constant S_{nh4+}		

Table 7: Operating parameters.

Symbol	Description	Symbol	Description
T	Temperature	q_{in}	Liquid flow rate in digester
V_{liq}	Liquid volume	V_{gas}	Gaseous volume
$S_{in},$	Inflow of component S or X	k_p	Gas outflow resistance
X_{in}			

Equations

From Equations (2) to (23), the reaction rates are given. The inhibition terms are given in Equations (24) to (31). The ODEs are given in Equations (35) to (78). The AEs are given in Equations (88) to (97).

The reaction rates ρ_i are the following:

$$\rho_1 = k_{dis}X_c \quad (2)$$

$$\rho_2 = k_{hyd,ch}X_{ch} \quad (3)$$

$$\rho_3 = k_{hyd,pr}X_{pr} \quad (4)$$

$$\rho_4 = k_{hyd,li}X_{li} \quad (5)$$

$$\rho_5 = k_{m,su} \frac{S_{su}}{K_{S,su} + S_{su}} X_{su} I_{IN,lim} I_{pH,aa} \quad (6)$$

$$\rho_6 = k_{m,aa} \frac{S_{aa}}{K_{S,aa} + S_{aa}} X_{aa} I_{IN,lim} I_{pH,aa} \quad (7)$$

$$\rho_7 = k_{m,fa} \frac{S_{fa}}{K_{S,fa} + S_{fa}} X_{fa} I_{IN,lim} I_{pH,aa} I_{h2,fa} \quad (8)$$

$$\rho_8 = k_{m,c4} \frac{S_{va}}{K_{S,c4} + S_{va}} X_{c4} \frac{S_{va}}{S_{bu} + S_{va}} I_{IN,lim} I_{pH,aa} I_{h2,c4} \quad (9)$$

$$\rho_9 = k_{m,c4} \frac{S_{bu}}{K_{S,c4} + S_{bu}} X_{c4} \frac{S_{bu}}{S_{bu} + S_{va}} I_{pH,aa} I_{h2,c4} \quad (10)$$

$$\rho_{10} = k_{m,pro} \frac{S_{pro}}{K_{S,pro} + S_{pro}} X_{pro} I_{IN,lim} I_{pH,aa} I_{h2,pro} \quad (11)$$

$$\rho_{11} = k_{m,ac} \frac{S_{ac}}{K_{S,ac} + S_{ac}} X_{ac} I_{IN,lim} I_{pH,ac} I_{nh3} \quad (12)$$

$$\rho_{12} = k_{m,h2} \frac{S_{h2}}{K_{S,h2} + S_{h2}} X_{h2} I_{IN,lim} I_{pH,h2} \quad (13)$$

$$\rho_{13} = k_{dec,Xsu}X_{su} \quad (14)$$

$$\rho_{14} = k_{dec,Xaa}X_{aa} \quad (15)$$

$$\rho_{15} = k_{dec,Xfa}X_{fa} \quad (16)$$

$$\rho_{16} = k_{dec,Xc4}X_{c4} \quad (17)$$

$$\rho_{17} = k_{dec,Xpro}X_{pro} \quad (18)$$

$$\rho_{18} = k_{dec,Xac}X_{ac} \quad (19)$$

$$\rho_{19} = k_{\text{dec}, \text{Xh2}} X_{\text{h2}} \quad (20)$$

$$\rho_{\text{T},8} = k_{\text{L}} a(S_{\text{h2}} - K_{\text{H},\text{h2}} RT S_{\text{gas},\text{h2}}) \quad (21)$$

$$\rho_{\text{T},9} = k_{\text{L}} a(S_{\text{ch4}} - K_{\text{H},\text{ch4}} RT S_{\text{gas},\text{ch4}}) \quad (22)$$

$$\rho_{\text{T},10} = k_{\text{L}} a(S_{\text{co2}} - K_{\text{H},\text{co2}} RT S_{\text{Xgas},\text{co2}}) \quad (23)$$

The inhibition terms are the following:

$$I_{\text{h2},\text{fa}} = \frac{K_{\text{I},\text{h2},\text{fa}}}{K_{\text{I},\text{h2},\text{fa}} + S_{\text{h2}}} \quad (24)$$

$$I_{\text{h2},\text{c4}} = \frac{K_{\text{I},\text{h2},\text{c4}}}{K_{\text{I},\text{h2},\text{c4}} + S_{\text{h2}}} \quad (25)$$

$$I_{\text{h2},\text{pro}} = \frac{K_{\text{I},\text{h2},\text{pro}}}{K_{\text{I},\text{h2},\text{pro}} + S_{\text{h2}}} \quad (26)$$

$$I_{\text{nh3}} = \frac{K_{\text{I},\text{nh3}}}{K_{\text{I},\text{nh3}} + S_{\text{nh3}}} \quad (27)$$

$$I_{\text{IN},\text{lim}} = \frac{S_{\text{IN}}}{S_{\text{IN}} + K_{\text{S},\text{IN}}} \quad (28)$$

$$I_{\text{pH},\text{aa}} = \frac{K_{\text{pH}}^{n_{\text{aa}}}}{S_{\text{h}^+}^{n_{\text{aa}}} + K_{\text{pH},\text{aa}}^{n_{\text{aa}}}} \quad (29)$$

$$I_{\text{pH},\text{ac}} = \frac{K_{\text{pH}}^{n_{\text{ac}}}}{S_{\text{h}^+}^{n_{\text{ac}}} + K_{\text{pH},\text{aa}}^{n_{\text{ac}}}} \quad (30)$$

$$I_{\text{pH},\text{h2}} = \frac{K_{\text{pH}}^{n_{\text{h2}}}}{S_{\text{h}^+}^{n_{\text{h2}}} + K_{\text{pH},\text{h2}}^{n_{\text{h2}}}} \quad (31)$$

with:

$$K_{\text{pH},\text{aa}} = 10^{\frac{\text{pH}_{\text{aa},\text{LL}} + \text{pH}_{\text{aa},\text{UL}}}{2}}, \quad n_{\text{pH},\text{aa}} = \frac{3}{\text{pH}_{\text{aa},\text{UL}} - \text{pH}_{\text{aa},\text{LL}}} \quad (32)$$

$$K_{\text{pH},\text{ac}} = 10^{\frac{\text{pH}_{\text{ac},\text{LL}} + \text{pH}_{\text{ac},\text{UL}}}{2}}, \quad n_{\text{pH},\text{ac}} = \frac{3}{\text{pH}_{\text{ac},\text{UL}} - \text{pH}_{\text{ac},\text{LL}}} \quad (33)$$

$$K_{\text{pH},\text{h2}} = 10^{\frac{\text{pH}_{\text{h2},\text{LL}} + \text{pH}_{\text{h2},\text{UL}}}{2}}, \quad n_{\text{pH},\text{h2}} = \frac{3}{\text{pH}_{\text{aa},\text{UL}} - \text{pH}_{\text{h2},\text{LL}}} \quad (34)$$

The following differential and algebraic equations make up the model:

$$\frac{dS_{\text{su}}}{dt} = \frac{q_{\text{in}}}{V_{\text{liq}}} (S_{\text{su},\text{in}} - S_{\text{su}}) + \rho_2 + (1 - f_{\text{fa},\text{li}}) \rho_4 - \rho_5 \quad (35)$$

$$\frac{dS_{\text{aa}}}{dt} = \frac{q_{\text{in}}}{V_{\text{liq}}} (S_{\text{aa},\text{in}} - S_{\text{aa}}) + \rho_3 - \rho_6 \quad (36)$$

$$\frac{dS_{\text{fa}}}{dt} = \frac{q_{\text{in}}}{V_{\text{liq}}} (S_{\text{fa},\text{in}} - S_{\text{fa}}) + f_{\text{fa},\text{li}} \rho_4 - \rho_7 \quad (37)$$

$$\frac{dS_{\text{va}}}{dt} = \frac{q_{\text{in}}}{V_{\text{liq}}} (S_{\text{va},\text{in}} - S_{\text{va}}) + (1 - Y_{\text{aa}}) f_{\text{va},\text{aa}} \rho_6 - \rho_8 \quad (38)$$

$$\frac{dS_{bu}}{dt} = \frac{q_{in}}{V_{liq}}(S_{bu,in} - S_{bu}) + (1 - Y_{su})f_{bu,su}\rho_5 \quad (39)$$

$$+ (1 - Y_{aa})f_{bu,aa}\rho_6 - \rho_9 \quad (40)$$

$$\begin{aligned} \frac{dS_{pro}}{dt} = \frac{q_{in}}{V_{liq}}(S_{pro,in} - S_{pro}) + (1 - Y_{su})f_{pro,su}\rho_5 \\ + (1 - Y_{aa})f_{pro,aa}\rho_6 + (1 - Y_{c4})0.54\rho_8 - \rho_{10} \end{aligned} \quad (41)$$

$$\begin{aligned} \frac{dS_{ac}}{dt} = \frac{q_{in}}{V_{liq}}(S_{ac,in} - S_{ac}) + (1 - Y_{su})f_{ac,su}\rho_5 \\ + (1 - Y_{aa})f_{ac,aa}\rho_6 + (1 - Y_{fa})0.7\rho_7 \\ + (1 - Y_{c4})0.31\rho_8 + (1 - Y_{c4})0.8\rho_9 \\ + (1 - Y_{pro})0.57\rho_{10} - \rho_{11} \end{aligned} \quad (42)$$

$$\begin{aligned} \frac{dS_{h2}}{dt} = \frac{q_{in}}{V_{liq}}(S_{h2,in} - S_{h2}) + (1 - Y_{su})f_{h2,su}\rho_5 \\ + (1 - Y_{aa})f_{h2,aa}\rho_6 + (1 - Y_{fa})0.3\rho_7 \\ + (1 - Y_{c4})0.15\rho_8 + (1 - Y_{c4})0.2\rho_9 \\ + (1 - Y_{pro})0.43\rho_{10} - \rho_{12} - \rho_{T,8} \end{aligned} \quad (43)$$

$$\begin{aligned} \frac{dS_{ch4}}{dt} = \frac{q_{in}}{V_{liq}}(S_{ch4,in} - S_{ch4}) + (1 - Y_{ac})\rho_{11} + (1 - Y_{h2})\rho_{12} \\ - \rho_{T,9} \end{aligned} \quad (44)$$

$$\begin{aligned} \frac{dS_{IC}}{dt} = \frac{q_{in}}{V_{liq}}(S_{IC,in} - S_{IC}) - \sum_{j=1}^{19} \left(\sum_{i=1-9,11-24} C_i v_{i,j} \rho_j \right) \\ - \rho_{T,10} \end{aligned} \quad (45)$$

$$\begin{aligned} \frac{dS_{IN}}{dt} = \frac{q_{in}}{V_{liq}}(S_{IN,in} - S_{IN}) - Y_{su}N_{bac}\rho_5 \\ + (N_{aa} - Y_{aa}N_{bac})\rho_6 - Y_{fa}N_{bac}\rho_7 \\ - Y_{c4}N_{bac}\rho_8 - Y_{c4}N_{bac}\rho_9 - Y_{pro}N_{bac}\rho_{10} \\ - Y_{ac}N_{bac}\rho_{11} - Y_{h2}N_{bac}\rho_{12} \end{aligned} \quad (46)$$

$$\begin{aligned} + (N_{bac} - N_{xc}) \sum_{i=13}^{19} \rho_i \\ + (N_{xc} - f_{xI,xc}N_I - f_{sI,xc}N_I - f_{pr,xc}N_{aa})\rho_1 \end{aligned}$$

$$\frac{dS_I}{dt} = \frac{q_{in}}{V_{liq}}(S_{I,in} - S_I) + f_{sI,xc}\rho_1 \quad (47)$$

$$\frac{dX_c}{dt} = \frac{q_{in}}{V_{liq}}(X_{c,in} - X_c) - \rho_1 + \sum_{i=13}^{19} \rho_i \quad (48)$$

$$\frac{dX_{ch}}{dt} = \frac{q_{in}}{V_{liq}}(X_{ch,in} - X_{ch}) + f_{ch,xc}\rho_1 - \rho_2 \quad (49)$$

$$\frac{dX_{pr}}{dt} = \frac{q_{in}}{V_{liq}}(X_{pr,in} - X_{pr}) + f_{pr,xc}\rho_1 - \rho_3 \quad (50)$$

$$\frac{dX_{li}}{dt} = \frac{q_{in}}{V_{liq}}(X_{li,in} - X_{li}) + f_{li,xc}\rho_1 - \rho_4 \quad (51)$$

$$\frac{dX_{su}}{dt} = \frac{q_{in}}{V_{liq}}(X_{su,in} - X_{su}) + Y_{su}\rho_5 - \rho_{13} \quad (52)$$

$$\frac{dX_{aa}}{dt} = \frac{q_{in}}{V_{liq}}(X_{aa,in} - X_{aa}) + Y_{aa}\rho_6 - \rho_{14} \quad (53)$$

$$\frac{dX_{fa}}{dt} = \frac{q_{in}}{V_{liq}}(X_{fa,in} - X_{fa}) + Y_{fa}\rho_7 - \rho_{15} \quad (54)$$

$$\frac{dX_{c4}}{dt} = \frac{q_{in}}{V_{liq}}(X_{c4,in} - X_{c4}) + Y_{c4}\rho_8 + Y_{c4}\rho_9 - \rho_{16} \quad (55)$$

$$\frac{dX_{pro}}{dt} = \frac{q_{in}}{V_{liq}}(X_{pro,in} - X_{pro}) + Y_{pro}\rho_{10} - \rho_{17} \quad (56)$$

$$\frac{dX_{ac}}{dt} = \frac{q_{in}}{V_{liq}}(X_{ac,in} - X_{ac}) + Y_{ac}\rho_{11} - \rho_{18} \quad (57)$$

$$\frac{dX_{h2}}{dt} = \frac{q_{in}}{V_{liq}}(X_{h2,in} - X_{h2}) + Y_{h2}\rho_{12} - \rho_{19} \quad (58)$$

$$\frac{dX_I}{dt} = \frac{q_{in}}{V_{liq}}(X_{I,in} - X_I) + f_{xI,xc}\rho_1 \quad (59)$$

$$\frac{dS_{cat+}}{dt} = \frac{q_{in}}{V_{liq}}(S_{cat+,in} - S_{cat+}) \quad (60)$$

$$\frac{dS_{an-}}{dt} = \frac{q_{in}}{V_{liq}}(S_{an-,in} - S_{an-}) \quad (61)$$

$$\frac{dS_{gas,h2}}{dt} = - \frac{S_{gas,h2}RTV_{liq}(\frac{\rho_{T,8}}{16} + \frac{\rho_{T,9}}{64} + \rho_{T,10})}{(p_{atm} - p_{gas,h2o})V_{gas}} + \rho_{T,8}\frac{V_{liq}}{V_{gas}} \quad (62)$$

$$\frac{dS_{gas,ch4}}{dt} = - \frac{S_{gas,ch4}RTV_{liq}(\frac{\rho_{T,8}}{16} + \frac{\rho_{T,9}}{64} + \rho_{T,10})}{(p_{atm} - p_{gas,h2o})V_{gas}} + \rho_{T,9}\frac{V_{liq}}{V_{gas}} \quad (63)$$

$$\frac{dS_{gas,co2}}{dt} = - \frac{S_{gas,co2}RTV_{liq}(\frac{\rho_{T,8}}{16} + \frac{\rho_{T,9}}{64} + \rho_{T,10})}{(p_{atm} - p_{gas,h2o})V_{gas}} + \rho_{T,10}\frac{V_{liq}}{V_{gas}} \quad (64)$$

The sum in Equation (45) is:

$$\begin{aligned} & \sum_{j=1}^{19} \left(\sum_{i=1-9,11-24} C_i v_{i,j} \rho_j \right) \\ &= \sum_{k=1}^{12} s_k \rho_k + s_{13}(\rho_{13} + \rho_{14} + \rho_{15} + \rho_{16} + \rho_{17} + \rho_{18} + \rho_{19}) \end{aligned} \quad (65)$$

where:

$$s_1 = -C_{xc} + f_{sI,xc}C_{sI} + f_{ch,xc}C_{ch} + f_{pr,xc}C_{pr} + f_{li,xc}C_{li}$$

$$+ f_{xI,xc} C_{xI} \quad (66)$$

$$s_2 = -C_{ch} + C_{su} \quad (67)$$

$$s_3 = -C_{pr} + C_{aa} \quad (68)$$

$$s_4 = -C_{li} + (1 - f_{fa,li}) C_{su} + f_{fa,li} C_{fa} \quad (69)$$

$$s_5 = -C_{su} + (1 - Y_{su})(f_{bu,su} C_{bu} + f_{pro,su} C_{pro} + f_{ac,su} C_{ac}) + Y_{su} C_{bac} \quad (70)$$

$$s_6 = -C_{aa} + (1 - Y_{aa})(f_{va,aa} C_{va} + f_{bu,aa} C_{bu} + f_{pro,aa} C_{pro} + f_{ac,aa} C_{ac}) + Y_{aa} C_{bac} \quad (71)$$

$$s_7 = -C_{fa} + (1 - Y_{fa}) 0.7 C_{ac} + Y_{fa} C_{bac} \quad (72)$$

$$s_8 = -C_{va} + (1 - Y_{c4}) 0.54 C_{pro} + (1 - Y_{c4}) 0.31 C_{ac} + Y_{c4} C_{bac} \quad (73)$$

$$s_9 = -C_{bu} + (1 - Y_{c4}) 0.8 C_{ac} + Y_{c4} C_{bac} \quad (74)$$

$$s_{10} = -C_{pro} + (1 - Y_{pro}) 0.57 C_{ac} + Y_{pro} C_{bac} \quad (75)$$

$$s_{11} = -C_{ac} + (1 - Y_{ac}) C_{ch4} + Y_{ac} C_{bac} \quad (76)$$

$$s_{12} = (1 - Y_{h2}) C_{ch4} + Y_{h2} C_{bac} \quad (77)$$

$$s_{13} = -C_{bac} + C_{xc} \quad (78)$$

The N-differences between the states are the following:

$$r_1 = N_{xc} - f_{xI,xc} N_I - f_{sI,xc} N_I - f_{pr,xc} N_{aa} \quad (79)$$

$$r_2 = -Y_{su} N_{bac} \quad (80)$$

$$r_3 = (N_{aa} - Y_{aa} N_{bac}) \quad (81)$$

$$r_4 = -Y_{fa} N_{bac} \quad (82)$$

$$r_5 = -Y_{c4} N_{bac} \quad (83)$$

$$r_6 = -Y_{pro} N_{bac} \quad (84)$$

$$r_7 = -Y_{ac} N_{bac} \quad (85)$$

$$r_8 = -Y_{h2} N_{bac} \quad (86)$$

$$r_9 = N_{bac} - N_{xc} \quad (87)$$

The 10 AEs for the DAE-implementation of ADM1.

$$0 = S_{cat+} + S_{nh4+} + S_{h+} - S_{hco3-} - \frac{S_{ac-}}{64} - \frac{S_{pro-}}{112} - \frac{S_{bu-}}{160} - \frac{S_{va-}}{208} - S_{oh-} - S_{an-} \quad (88)$$

$$0 = S_{oh-} - \frac{K_w}{S_{h+}} \quad (89)$$

$$0 = S_{va-} - \frac{K_{a,va} S_{va}}{K_{a,va} + S_{h+}} \quad (90)$$

$$0 = S_{bu-} - \frac{K_{a,bu} S_{bu}}{K_{a,bu} + S_{h+}} \quad (91)$$

$$0 = S_{pro-} - \frac{K_{a,pro} S_{pro}}{K_{a,pro} + S_{h+}} \quad (92)$$

$$0 = S_{ac^-} - \frac{K_{a,ac}S_{ac}}{K_{a,ac} + S_{h^+}} \quad (93)$$

$$0 = S_{hco3^-} - \frac{K_{a,co2}S_{IC}}{K_{a,co2} + S_{h^+}} \quad (94)$$

$$0 = S_{nh4^+} - \frac{S_{h^+}S_{IN}}{K_{a,IN} + S_{h^+}} \quad (95)$$

$$0 = S_{IC} - S_{co2} - S_{hco3^-} \quad (96)$$

$$0 = S_{IN} - S_{nh3} - S_{nh4^+} \quad (97)$$

The gaseous outflow:

$$q_{gas} = k_p(p_{gas} - P_{atm}) \quad (98)$$

with

$$p_{gas} = S_{gas,h2} \frac{RT}{16} + S_{gas,ch4} \frac{RT}{64} + S_{gas,co2} RT + p_{gas,h2o} \quad (99)$$

Appendix B. Numerical values of states and parameters

Over the course of this work numerical values are substituted for states, e.g., as initial conditions and parameters. Unless otherwise stated, the numerical values presented in this appendix are used.

Table 1: Steady-state state variables with $q_{liq} = 170.0 \text{ m}^3\text{d}^{-1}$, calculated by applying the input variables of Table 3 for 1000 days.

State	Value	State	Value
S_{su}	0.01195	X_{c4}	0.43192
S_{aa}	0.00531	X_{pro}	0.13731
S_{fa}	0.09862	X_{ac}	0.76056
S_{va}	0.01162	X_{h2}	0.31702
S_{bu}	0.01325	X_I	25.61740
S_{pro}	0.01578	S_{cat+}	0.04
S_{ac}	0.19762	S_{an-}	0.02
S_{h2}	2.3595E-7	S_{h+}	3.4234E-8
S_{ch4}	0.05509	S_{va}	0.011560
S_{IC}	0.15268	S_{bu}	0.01322
S_{IN}	0.13023	S_{pro}	0.01574
S_I	0.32870	S_{ac}	0.19724
X_{xc}	0.30869	S_{hco3-}	0.14278
X_{ch}	0.02795	S_{co2}	0.00990
X_{pr}	0.10258	S_{nh3}	0.00409
X_{li}	0.02948	S_{nh4+}	0.12613
X_{su}	0.42017	$S_{gas,h2}$	1.0241E-5
X_{aa}	1.17918	$S_{gas,ch4}$	1.6256
X_{fa}	0.24303	$S_{gas,co2}$	0.014151

Table 2: Steady-state state variables with $q_{liq} = 85.0 \text{ m}^3\text{d}^{-1}$, calculated by applying the input variables of Table 3 for 1000 days.

State	Value	State	Value
S_{su}	0.00762	X_{pro}	0.10899
S_{aa}	0.0033953	X_{ac}	0.60874
S_{fa}	0.05728	X_{h2}	0.25345
S_{va}	0.0072466	X_I	25.804
S_{bu}	0.0082609	S_{cat+}	0.04
S_{pro}	0.0093745	S_{an-}	0.02
S_{ac}	0.095295	S_{h+}	3.2168E-8
S_{h2}	1.4997E-7	S_{oh-}	6.4622E-7

S_{ch4}	0.050567	S_{va}	0.0072298
S_{IC}	0.15725	S_{bu}	0.0082434
S_{IN}	0.13375	S_{pro}	0.0093517
S_I	0.42216	S_{ac}	0.095119
X_{xc}	0.20108	S_{hco3-}	0.14763
X_{ch}	0.014475	S_{co2}	0.009619
X_{pr}	0.051881	S_{nh3}	0.0044624
X_{li}	0.015478	S_{nh4+}	0.12929
X_{su}	0.33899	$S_{gas,h2}$	7.1083E-6
X_{aa}	0.92778	$S_{gas,ch4}$	1.5829
X_{fa}	0.19973	$S_{gas,co2}$	0.013788
X_{c4}	0.34071		

Table 3: Input variables.

Input	Value	Input	Value		
$S_{su,in}$	0.01	$S_{IC,in}$	0.04	$X_{aa,in}$	0.01
$S_{aa,in}$	0.001	$S_{IN,in}$	0.01	$X_{fa,in}$	0.01
$S_{fa,in}$	0.001	$S_{I,in}$	0.02	$X_{c4,in}$	0.01
$S_{va,in}$	0.001	$S_{cat,in}$	0.04	$X_{pro,in}$	0.01
$S_{bu,in}$	0.001	$S_{an,in}$	0.02	$X_{ac,in}$	0.01
$S_{pro,in}$	0.001	$X_{ch,in}$	5.0	$X_{h2,in}$	0.01
$S_{ac,in}$	0.001	$X_{pr,in}$	20.0	$X_{I,in}$	25.0
$S_{h2,in}$	1.0E-8	$X_{li,in}$	5.0	$X_{xc,in}$	2.0
$S_{ch4,in}$	1.0E-5	$X_{su,in}$	0.0		

Table 4: Numerical values stoichiometric parameters.

Parameter	Value	Parameter	Value		
$f_{sI,xc}$	0.1	$f_{bu,su}$	0.13	Y_{pro}	0.04
$f_{xI,xc}$	0.2	$f_{va,aa}$	0.23	Y_{aa}	0.08
$f_{ch,xc}$	0.2	$f_{va,aa}$	0.23	Y_{c4}	0.06
$f_{pr,xc}$	0.2	$f_{h2,aa}$	0.06	Y_{fa}	0.06
$f_{li,xc}$	0.3	$f_{ac,aa}$	0.40	Y_{h2}	0.06
$f_{ac,su}$	0.41	$f_{bu,aa}$	0.26	Y_{ac}	0.05
$f_{pro,su}$	0.27	$f_{fa,li}$	0.95		
$f_{h2,su}$	0.19	Y_{su}	0.1		

Table 5: Numerical value composition parameters.

Parameter	Value	Parameter	Value
N_{xc}	0.0027	C_{su}	0.0313
N_I	0.0043	C_{aa}	0.03
N_{aa}	0.007	C_{fa}	0.0217
N_{bac}	0.0057	C_{bac}	0.0313
C_{xc}	0.027863	C_{bu}	0.025
C_{sI}	0.03	C_{pro}	0.0268
C_{ch}	0.0313	C_{ac}	0.0313
C_{pr}	0.03	C_{va}	0.024
C_{li}	0.022	C_{ch4}	0.0156
C_{xI}	0.03		

Table 6: Numerical values kinetic parameters.

Parameter	Value	Parameter	Value
k_{dis}	0.5	$K_{I,nh3}$	0.0018
$k_{hyd,ch}$	10	$K_{I,h2,pro}$	3.5E-6
$k_{hyd,pr}$	10	$K_{I,h2,c4}$	1E-5
$k_{hyd,li}$	10	$K_{I,h2,fa}$	5E-6
$k_{m,su}$	30	$K_{S,IN}$	1E-4
$K_{S,su}$	0.5	$k_{dec,Xaa}$	0.02
$k_{m,aa}$	50	$k_{dec,Xc4}$	0.02
$K_{S,aa}$	0.3	$k_{dec,Xfa}$	0.02
$k_{m,fa}$	6	$k_{dec,Xpro}$	0.02
$K_{S,fa}$	0.4	$k_{dec,Xac}$	0.02
$k_{m,c4}$	20	$k_{dec,Xh2}$	0.02
$K_{S,c4}$	0.2	$k_{dec,Xsu}$	0.02
$k_{m,pro}$	13	$pH_{LL,aa}$	5.5
$K_{S,pro}$	0.1	$pH_{UL,aa}$	4
$k_{m,ac}$	8	$pH_{LL,h2}$	6
$K_{S,ac}$	0.15	$pH_{UL,h2}$	5
$k_{m,h2}$	35	$pH_{LL,ac}$	7
$K_{S,h2}$	7E-6	$pH_{UL,ac}$	6
$k_L a$	200		

Table 7: Numerical values physico-chemical constants.

Parameter	Value	Unit	Parameter	Value	Unit
R	8.3145e-2	bar M ⁻¹ K ⁻¹	$K_{a,co2}$	4.9371E-7	M
K_w	2.0788E-14	M	$K_{a,IN}$	1.1110E-9	M
$K_{a,va}$	1.3804E-5	M	$K_{H,co2}$	2.7147E-2	M _{liq} bar ⁻¹
$K_{a,bu}$	1.5136E-5	M	$K_{H,h2}$	7.3847E-4	M _{liq} bar ⁻¹
$K_{a,pro}$	1.3183E-5	M	$K_{H,ch4}$	1.1619E-3	M _{liq} bar ⁻¹

$$K_{a,ac} \quad 1.7378E-5 \quad M \quad |$$

Table 8: Numerical values operating parameters.

Parameter	Value	Unit	Parameter	Value	Unit
V_{liq}	3400	m^3	$p_{gas,h2o}$	0.0313	bar
V_{gas}	300	m^3	T	308.15	K
P_{atm}	1.013	bar	k_p	5E4	$m^3 \text{ d}^{-1} \text{ bar}^{-1}$

Appendix C. Relations between ADM1 model structure and structural identifiability results

Interactions

Following interactions are found in the model structure of ADM1:

- Through the different degradation processes ρ_i , $i = 1, \dots, 19$ each state directly influences the concentration of its degradation products, and for $\rho_5 - \rho_{12}$ the biomass performing this degradation.
- Each biomass (X_{su} , X_{aa} , X_{fa} , X_{pro} , X_{c4} , X_{h2} , X_{ac}) influences the concentration of the component it degrades and its degradation products.
- All biomass decays to X_{xc} which is the component at the beginning of the reaction network. In that way, each biomass has an indirect influence on *every* component in the reaction scheme.
- For each reaction, there is an interaction with the source/sink components S_{IC} and S_{IN} to compensate for an excess/deficiency of carbon and nitrogen between the component and its degradation products. The flows from/to S_{IC} and S_{IN} are indicated with s_i , $i = 1, \dots, 1$, r_i , $i = 1, \dots, 9$, respectively on Figure 1. Thus, S_{IC} and S_{IN} are influenced by almost all reactions.
- Inorganic nitrogen S_{IN} directly influences the acetate degradation since its ammonia fraction S_{nh3} is inhibiting. If nitrogen is a limiting component in biomass growth, it also influences the reactions performed by microbial groups, i.e., $\rho_5 - \rho_{12}$.
- There are three gaseous components $S_{gas,co2}$, $S_{gas,h2}$ and $S_{gas,ch4}$. The outflow of each of these gases q_{gas} out of the system is determined by the quantity of *all* of each of these gases, thus, they influence each other.
- Each of the gases exchanges mass with its soluble form, i.e., S_{h2} , S_{ch4} , S_{co2} . Thus, the soluble forms, through the gaseous form, interact with each other. S_{co2} is a pH-determined fraction of the inorganic carbon S_{IC} , thus through S_{co2} and $S_{gas,co2}$, S_{IC} has an influence on S_{h2} , S_{ch4} , S_{co2} .
- Valeric acid (S_{va}), butyric acid (S_{bu}), acetic acid (S_{ac}), propionic acid (S_{pro}), inorganic nitrogen (S_{IN}) and inorganic carbon (S_{IC}) directly influence the pH.
- The pH has a direct inhibition effect on reactions $\rho_5 - \rho_{12}$, and determines the fraction of S_{IC} and S_{IN} existing as S_{co2} and S_{nh3} , respectively.
- Hydrogen S_{h2} inhibits the degradation of S_{bu} , S_{va} , S_{fa} and S_{pro} .

Identifiability and interconnectivity

To decrease the connectivity, three hypothetical ‘storage’ states were introduced γ_1 , γ_2 and γ_3 . γ_1 collects all COD from decayed biomass and γ_2 and γ_3 collect the excess of C and N originating from the decay of biomass. In this manner, the COD, C and N is not recirculated to the particulate composites X_{xc} via the ‘decay’ recirculation (see Figure 1).

The observability results of this ‘modified’ reaction network are then the following. When X_{xc} is measured, only $X_{xc}(0)$ is observable. When only S_{su} is measured, only $X_{xc}(0)$, $X_{ch}(0)$ and $S_{su}(0)$ are observable. Thus, only information from the states themselves or of states ‘upstream’ in the reaction pathway can be observed. Because of the removal of the feedback via the decay, there is no information pathway linking $X_{xc}(0)$ or S_{su} to the rest of the system. When one goes further downstream in the altered reaction network, the observability results become applicable again. For instance when only S_{pro} is measured almost all states and parameters are observable with the exception of the parameters and initial conditions already listed in Table 1: $f_{sI,xc}$, $f_{xI,xc}$, C_{sI} , C_{xI} , $S_I(0)$, $X_I(0)$, S_{cat+} , S_{an-} , and $\gamma_1(0)$, $\gamma_2(0)$, $\gamma_3(0)$ and the composition parameters N_I , C_{xc} and N_{xc} . The composition parameters become non-observable because of the carbon and nitrogen captured in the non-observable $\gamma_2(0)$ and $\gamma_3(0)$. Since S_{pro} is connected to the rest of the system via multiple mechanisms, measuring it is thus sufficiently informative from an observability or identifiability point of view. Finally, note that it was necessary to introduce *all* ‘storage’ states γ_1 , γ_2 and γ_3 to break open the connectivity. One or two of them was insufficient.

Table 1: The results of the parameters of ADM1 for the scenarios with inputs q_{liq} and/or $S_{x,in}$. Observability is equivalent to structurally locally identifiable.

Measurement	Result
All states	All parameters observable excl. C_{sI} , C_{xI}
Any state excl. S_I , X_I , S_{cat+} , S_{an-}	All parameters observable excl. $f_{sI,xc}$, $f_{xI,xc}$, C_{sI} , C_{xI} , $S_I(0)$, $X_I(0)$, $S_{cat+}(0)$, $S_{an-}(0)$
S_I	All parameters observable excl. C_{sI} , C_{xI} , $X_I(0)$, $S_{cat+}(0)$, $S_{an-}(0)$
X_I	All parameters observable excl. C_{sI} , C_{xI} , $S_I(0)$, $S_{cat+}(0)$, $S_{an-}(0)$
S_{cat+}	None observable excl. $S_{cat+}(0)$
S_{an-}	None observable excl. $S_{an-}(0)$

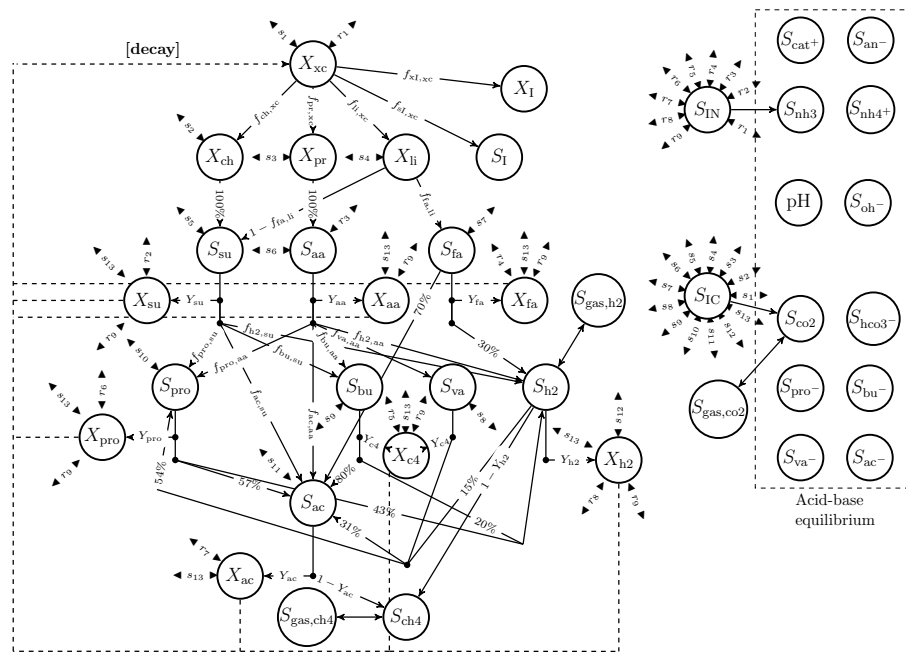


Figure 1: **The structure of ADM1, on which the stoichiometric parameters $f_{i,j}$ and Y_i are indicated.** Inorganic nitrogen S_{IC} and carbon S_{IN} are source/sink components that collect or make up excess or deficiencies of carbon and nitrogen, due to different N- or C-compositions of the different components in the system, indicated with r_i and s_i respectively. A separate system defines the acid-base equilibrium. The microbial metabolism on the substrates is inhibited by the pH-level, limiting inorganic nitrogen, ammonia and hydrogen.

Appendix D. In-depth discussion of the state profiles in Experiment 2

The state profiles for Experiment 2 are shown in Figure 1. Notable are the rises in the particulates X_{xc} , X_{ch} , X_{pr} and X_{li} due to the change of liquid inflow at $t = 25$ d and the peaks in S_{su} and S_{aa} due to the elevated inflow and lowered pH. Note that the lowering of the pH also inhibits the degradation of S_{fa} , S_{va} , S_{bu} , S_{pro} and S_{h2} allowing the concentrations of these state to rise dramatically. For S_{va} , S_{bu} , S_{pro} , the fermentation of the added S_{su} and S_{aa} gives an additional rise. After reactivation of the biomass by raising the pH to a normal level, the concentrations of the mentioned states drop significantly. The concentration of S_{ac} also increases due to the biomass inhibition, but increases even more after reactivation of the biomass. This is because S_{ac} is the product of the degradation of S_{va} , S_{bu} , S_{pro} .

The concentrations of biomass (X_{su} , X_{aa} , X_{fa} , X_{c4} , X_{pro} , X_{ac} and X_{h2}) decrease during the period of low pH. Important to note is the drop in inorganic carbon S_{IC} when the pH drops. In the acidification, a large portion of the inorganic carbon effervesces to gas, i.e., $S_{gas,co2}$ and leaves the CSTR in the gas outflow. When the digestion ‘re-activates’ when the pH is raised again, the C-consuming reaction also re-activates, further decreasing the C-content. However, since the inorganic carbon content is low, the limiting C-inhibition becomes active. The effect of this artificially added inhibition is negligible for almost all the states, i.e., the profiles with or without this inhibition are visually almost indiscernible. Only, for the lipids X_{li} a noticeable peak around $t = 60$ days can be observed, because of the inhibition of lipid uptake.

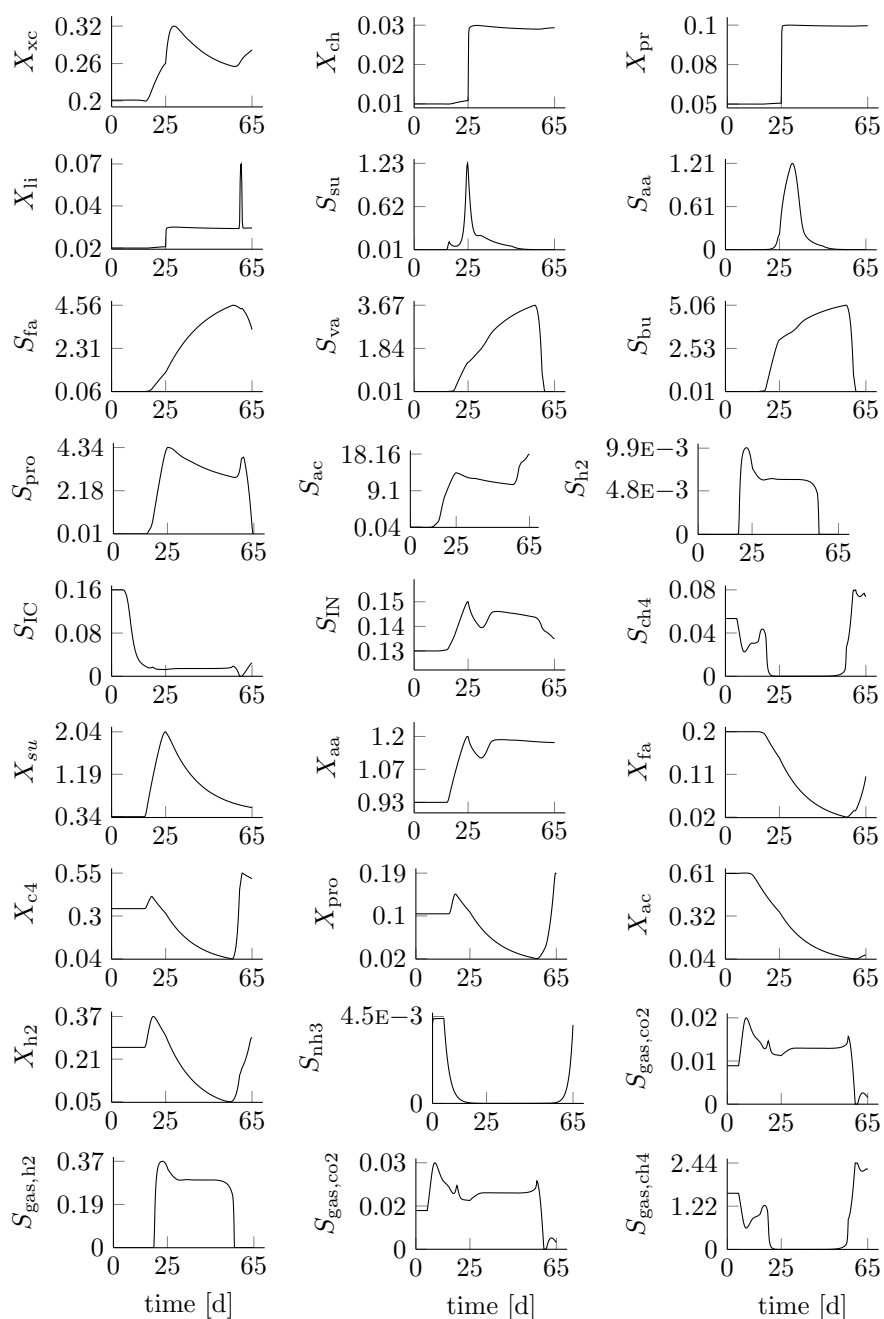


Figure 1: Concentration profiles of most states in Experiment 2.

Appendix E. Parameter estimation results

This appendix reports the parameter estimation results for the two experiments at different noise levels.

Experiment 1

The degradation of the states S_{su} , S_{aa} , S_{pro} , S_{fa} , S_{bu} , S_{va} , S_{h2} , S_{ac} by biomass is described by Monod kinetics and inhibited by H_2 , pH and NH_3 . The fact that the Monod parameter estimates are strongly interrelated and are not estimated well is not entirely surprising. No accurate results could have been expected if the values of the associated states are not below and above the K_S reference value. Because of the larger spread of acetic acid S_{ac} on the kinetic profiles, as shown in Figure 1, it is expected that both maximum uptake rate $k_{m,ac}$ (54.) and saturation constant $K_{S,ac}$ (55.) show accurate estimations.

Surprisingly, this is *only* so for the saturation constant and *not* for the maximum uptake rate. The reason for this poor estimation is most likely the strong negative interdependence with the NH_3 inhibition parameter $K_{I,nh3}$ (56.). This parameter appears indeed together with $k_{m,ac}$ in the equations for ρ_{11} . Since the NH_3 -values are generally very low and do not change much, $K_{I,nh3}$ is expected to be inaccurately estimated, and due to the exhibited interdependence with $k_{m,ac}$, the latter will also be estimated poorly. A similar reasoning can be made to explain the interdependence between (i) the Monod parameters of the degradation of S_{fa} (45.-46.) and the parameter related to its inhibition (47.), (ii) the Monod parameters of the degradation of S_{bu} , S_{va} (48.-49.) and the parameters related to its inhibition (50.), and (iii) the Monod parameters of the degradation of S_{pro} (51.-52.) and the parameter related to its inhibition (53.). Besides these interdependence between the Monod parameters and its inhibition constants, there is also a strong interdependence between the different sets of Monod parameters (42.-43.), (43.-44.), (45.-46.), (48.-49.), (51.-52.) and (54.) and the associated stoichiometric parameters (14.-22.). This interdependence is quite understandable, since there are multiple reactions pathways interconnecting the states S_{su} , S_{aa} , S_{va} , S_{pro} , S_{fa} , S_{bu} , S_{h2} , S_{ac} . Any change in the value of state induced by a bias in the estimate of a parameter, can then approximately be corrected by a change in value of another parameter in the network. Consider for instance the degradation of S_{su} to S_{pro} . Imagine that for some reason the value of $k_{m,su}$ is estimated too high. There are several ways to ‘correct’ this: the saturation constant $K_{S,su}$ could be increased, the stoichiometric parameter $f_{pro,su}$ can be decreased or the kinetic parameter $k_{m,pro}$ can be increased to alleviate the increase in the degradation product S_{pro} .

Finally, the connection between the composition parameters (23.-35.) and many of the stoichiometric parameters (10.-22.) and kinetic parameters (41.-58.) is explained. The fact that the C and N composition parameters are strongly connected to each other and the stoichiometric parameters most likely stem from the fact that they are primarily observed by the measurement of a single variable, i.e., the inorganic carbon S_{IC} or the inorganic nitrogen S_{IN} . A bias

in the estimate of one of the composition parameters can then approximately be offset by a change in any other composition parameters, or, by an off-set in the stoichiometric parameters. Since the stoichiometric parameters (10.-22.) are strongly related to the kinetic parameters (41.58.), a strong interdependence between the composition parameters and the stoichiometric parameters implies also a strong interdependence between the composition parameters and the kinetic parameters.

Table 1: Parameter estimation results of Experiment 1 for Gaussian measurement noise with a magnitude of 5% and 10% of the initial conditions. The true parameter values (θ_i^*), the relative deviation of the Monte Carlo estimate $\hat{\theta}_{MC,i\theta}$ and the relative standard deviation of the Monte Carlo estimate $\sigma_{MC,i\theta}$ are shown. ^(†) $C_I = f_{sI,xc}C_{sI} + f_{xI,xc}C_{xI}$. A bias or standard deviation of less than 5%, between 5% and 25% and more than 25% is indicated in white, light gray and dark gray respectively.

		5%		10%	
$\theta_{i\theta}$	$\theta_{i\theta}^*$	Bias _{rel,$i\theta$}	$\sigma_{rel,i\theta}$	Bias _{rel,$i\theta$}	$\sigma_{rel,i\theta}$
Stoichiometric parameters					
1. $f_{fa,li}$	0.95	2.6%	10.3%	−0.6%	19.2%
2. Y_{ac}	0.05	−1.2%	8.2%	−2.8%	12.2%
3. Y_{h2}	0.06	4.4%	15.9%	3.6%	25.6%
4. Y_{pro}	0.04	−0.6%	16.3%	−2.0%	26.5%
5. Y_{c4}	0.06	−3.2%	15.9%	−4.3%	26.7%
6. Y_{fa}	0.06	−3.9%	16.1%	−5.0%	27.5%
7. Y_{su}	0.1	1.7%	13.6%	−6.9%	37.2%
8. Y_{aa}	0.08	0.8%	6.6%	−2.8%	13.9%
9. $f_{sI,xc}$	0.1	−1.5%	5.5%	1.1%	8.6%
10. $f_{ch,xc}$	0.2	−3.3%	34.4%	−12.4%	64.9%
11. $f_{pr,xc}$	0.2	9.3%	50.2%	8.1%	76.5%
12. $f_{ii,xc}$	0.3	−0.2%	23.7%	5.1%	42.0%
13. $f_{xI,xc}$	0.2	−4.8%	24.4%	−4.0%	50.5%
14. $f_{bu,su}$	0.13	−54.8%	152.1%	−48.6%	154.2%
15. $f_{pro,su}$	0.27	−5.3%	78.5%	1.3%	83.7%
16. $f_{ac,su}$	0.41	32.9%	55.4%	23.1%	64.9%
17. $f_{h2,su}$	0.19	−26.0%	104.6%	−18.6%	114.6%
18. $f_{va,aa}$	0.23	1.6%	15.0%	−1.0%	23.2%
19. $f_{bu,aa}$	0.26	10.3%	20.9%	6.4%	29.3%
20. $f_{pro,aa}$	0.05	1.4%	78.8%	−19.6%	100.7%
21. $f_{ac,aa}$	0.4	−7.1%	19.0%	1.7%	29.2%
22. $f_{h2,aa}$	0.06	−5.0%	83.2%	−18.7%	115.1%
23. C_{xc}	0.02786	−193.9%	151.4%	−192.4%	132.4%
24. N_{xc}	0.00269	−98.8%	182.2%	−79.2%	174.4%
25. C_{bac}	0.0313	−162.7%	141.8%	−175.2%	121.5%
26. N_{bac}	0.0057	−43.3%	128.7%	−79.4%	155.5%
27. N_I	0.0043	−162.6%	178.4%	−161.1%	170.8%
28. $C_I^{(\dagger)}$	0.009	−120.6%	21.2%	−108.7%	19.7%
29. N_{aa}	0.007	−7.5%	22.3%	−13.4%	23.7%
30. C_{ch}	0.0313	−48.5%	154.5%	−65.5%	160.1%
31. C_{su}	0.0313	−76.5%	162.6%	−78.9%	173.3%
32. C_{pr}	0.03	−13.6%	63.8%	−16.3%	64.7%
33. C_{aa}	0.03	−78.9%	182.4%	−68.6%	171.4%
34. C_{li}	0.022	−63.2%	153.2%	−42.8%	149.7%
35. C_{fa}	0.0217	−61.0%	98.2%	−61.7%	106.6%

Kinetic parameters					
36. $k_L a$	200	0.1%	1.8%	0.7%	4.9%
37. k_{dis}	0.5	1.3%	5.2%	-1.6%	9.0%
38. $k_{hyd,ch}$	10	-0.2%	3.6%	-1.5%	7.0%
39. $k_{hyd,pr}$	10	0.3%	1.5%	0.2%	2.4%
40. $k_{hyd,li}$	10	0.2%	3.5%	0.5%	6.7%
41. $k_{m,su}$	30	-111.0%	180.3%	-140.7%	165.5%
42. $K_{S,su}$	0.5	-100.9%	171.1%	-139.4%	162.7%
43. $k_{m,aa}$	50	-113.0%	180.8%	-114.5%	177.0%
44. $K_{S,aa}$	0.3	-110.8%	181.1%	-113.2%	178.1%
45. $k_{m,fa}$	6	-39.7%	76.5%	-61.4%	96.2%
46. $K_{S,fa}$	0.4	-47.8%	96.3%	-73.2%	126.2%
47. $K_{I,h2,fa}$	5E-6	-138.7%	190.4%	-93.7%	180.3%
48. $k_{m,c4}$	20	-119.3%	139.2%	-110.2%	134.4%
49. $K_{S,c4}$	0.2	-130.8%	153.9%	-123.7%	153.5%
50. $K_{I,h2,c4}$	1E-5	-130.5%	185.8%	-154.9%	173.5%
51. $k_{m,pro}$	13	-99.8%	141.4%	-100.7%	120.1%
52. $K_{S,pro}$	0.1	-92.0%	136.6%	-101.0%	130.9%
53. $K_{I,h2,pro}$	3.5E-6	-115.8%	188.4%	-111.0%	179.1%
54. $k_{m,ac}$	8	-113.0%	128.9%	-124.5%	125.9%
55. $K_{S,ac}$	0.15	-0.7%	7.8%	-3.7%	14.4%
56. $K_{I,nh3}$	0.0018	9.7%	88.7%	-11.0%	133.3%
57. $k_{m,h2}$	35	-124.0%	163.8%	-91.8%	160.7%
58. $K_{S,h2}$	7E-5	-93.2%	150.8%	-57.3%	146.8%
59. $k_{dec,Xsu}$	0.02	-0.3%	35.2%	-8.7%	61.3%
60. $k_{dec,Xaa}$	0.02	3.7%	25.8%	-9.8%	53.2%
61. $k_{dec,Xfa}$	0.02	1.1%	32.7%	-3.1%	57.6%
62. $k_{dec,Xc4}$	0.02	2.8%	20.1%	2.4%	34.0%
63. $k_{dec,Xpro}$	0.02	1.9%	23.4%	-0.5%	41.4%
64. $k_{dec,Xac}$	0.02	2.1%	25.5%	1.5%	37.4%
65. $k_{dec,Xh2}$	0.02	7.4%	29.5%	3.7%	50.5%
66. $pH_{UL,aa}$	5.5	-18.0%	14.3%	-13.7%	15.3%
67. $pH_{LL,aa}$	4	-35.0%	12.8%	-34.3%	16.3%
68. $pH_{UL,h2}$	7	4.5%	11.1%	6.5%	12.8%
69. $pH_{LL,h2}$	6	6.4%	6.8%	7.2%	7.9%
70. $pH_{UL,ac}$	6	-15.5%	16.2%	-10.9%	18.8%
71. $pH_{LL,ac}$	5	-13.5%	11.2%	-12.4%	12.2%
Initial conditions					
72. $S_{su}(0)$	0.00762	0.4%	4.8%	0.1%	10.5%
73. $S_{aa}(0)$	0.00340	-0.9%	5.3%	0.0%	9.7%
74. $S_{fa}(0)$	0.0573	-0.1%	4.0%	-0.2%	9.6%
75. $S_{va}(0)$	0.00725	0.2%	3.2%	-1.0%	7.8%
76. $S_{bu}(0)$	0.00826	-0.3%	3.7%	-0.7%	6.9%
77. $S_{pro}(0)$	0.00937	-0.8%	3.7%	0.4%	7.1%
78. $S_{ac}(0)$	0.09540	0.3%	3.6%	0.1%	6.3%
79. $S_{h2}(0)$	1.500E-7	0.4%	4.8%	-0.5%	10.2%
80. $S_{ch4}(0)$	0.0505	0.5%	4.3%	0.5%	9.0%
81. $S_{IC}(0)$	0.157	0.0%	0.5%	-0.1%	0.9%
82. $S_{IN}(0)$	0.1337	0.0%	0.5%	-0.1%	1.0%
83. $S_I(0)$	0.422	-0.1%	0.9%	-0.2%	1.8%
84. $X_{xc}(0)$	0.201	0.2%	2.4%	0.5%	4.7%
85. $X_{ch}(0)$	0.0145	0.5%	4.9%	0.1%	10.3%
86. $X_{pr}(0)$	0.0520	-0.7%	4.8%	0.0%	9.6%
87. $X_{li}(0)$	0.0155	-0.7%	4.8%	0.1%	10.0%

88. $X_{su}(0)$	0.339	-0.1%	0.9%	0.2%	1.6%
89. $X_{aa}(0)$	0.928	-0.3%	0.9%	-0.2%	1.7%
90. $X_{fa}(0)$	0.200	0.0%	1.0%	-0.4%	1.9%
91. $X_{c4}(0)$	0.341	0.0%	0.9%	-0.2%	1.7%
92. $X_{pro}(0)$	0.109	-0.1%	1.0%	0.0%	1.9%
93. $X_{ac}(0)$	0.608	-0.1%	0.8%	0.1%	1.6%
94. $X_{h2}(0)$	0.253	-0.2%	0.9%	-0.5%	1.6%
95. $X_I(0)$	25.8	0.1%	1.0%	0.2%	2.0%
96. $S_{cat+}(0)$	0.04	-0.1%	0.8%	0.1%	1.5%
97. $S_{an-}(0)$	0.02	0.0%	0.8%	0.0%	1.5%
98. $S_{gas,h2}(0)$	7.108E-6	-0.3%	5.0%	1.6%	9.9%
99. $S_{gas,ch4}(0)$	1.582	0.2%	4.9%	-1.1%	9.8%
100. $S_{gas,co2}(0)$	0.0138	-0.1%	5.2%	-0.1%	11.2%

Table 2: Parameter estimation results of Experiment 1 for Gaussian measurement noise with a magnitude of 15% and 25% of the initial conditions. The true parameter values (θ_i^*), the relative deviation of the Monte Carlo estimate $\hat{\theta}_{MC,i\theta}$ and the relative standard deviation of the Monte Carlo estimate $\sigma_{MC,i\theta}$ are indicated. $(\dagger)C_I = f_{sI,xc}C_{sI} + f_{xI,xc}C_{xI}$. A bias or standard deviation of less than 5%, between 5% and 25% and more than 25% is indicated in white, light gray and dark gray respectively.

		15%		25%	
$\theta_{i\theta}$	$\theta_{i\theta}^*$	Bias _{rel,$i\theta$}	$\sigma_{rel,i\theta}$	Bias _{rel,$i\theta$}	$\sigma_{rel,i\theta}$
Stoichiometric parameters					
1. $f_{fa,li}$	0.95	−4.43%	29.4%	−6.16%	45.1%
2. Y_{ac}	0.05	−4.54%	18.3%	−16.22%	34.8%
3. Y_{h2}	0.06	4.89%	34.2%	−5.64%	52.6%
4. Y_{pro}	0.04	−10.50%	39.1%	−13.12%	60.3%
5. Y_{c4}	0.06	−10.38%	36.2%	−26.12%	59.8%
6. Y_{fa}	0.06	−7.73%	41.0%	−33.53%	79.8%
7. Y_{su}	0.1	−25.61%	73.7%	−60.57%	109.2%
8. Y_{aa}	0.08	−7.77%	21.1%	−18.16%	34.5%
9. $f_{sI,xc}$	0.1	3.71%	13.1%	13.22%	18.3%
10. $f_{ch,xc}$	0.2	−9.57%	74.7%	−21.90%	105.9%
11. $f_{pr,xc}$	0.2	6.03%	84.7%	7.04%	100.2%
12. $f_{li,xc}$	0.3	5.52%	54.1%	11.63%	67.0%
13. $f_{xI,xc}$	0.2	−6.59%	75.3%	−9.20%	91.3%
14. $f_{bu,su}$	0.13	−51.97%	159.2%	−17.48%	146.2%
15. $f_{pro,su}$	0.27	1.09%	88.1%	−3.33%	99.5%
16. $f_{ac,su}$	0.41	18.99%	65.8%	17.64%	71.3%
17. $f_{h2,su}$	0.19	−6.97%	122.1%	−21.36%	142.0%
18. $f_{va,aa}$	0.23	−0.13%	28.4%	1.89%	30.4%
19. $f_{bu,aa}$	0.26	7.03%	36.1%	5.91%	41.3%
20. $f_{pro,aa}$	0.05	−18.71%	112.8%	−56.26%	148.5%
21. $f_{ac,aa}$	0.4	7.32%	37.3%	12.15%	45.7%
22. $f_{h2,aa}$	0.06	−63.16%	145.9%	−67.00%	159.9%
23. C_{xc}	0.02786	−204.14%	120.1%	−165.36%	144.0%
24. N_{xc}	0.00269	−71.49%	173.4%	−94.76%	183.5%
25. C_{bac}	0.0313	−160.57%	129.6%	−170.15%	136.1%
26. N_{bac}	0.0057	−64.65%	160.5%	−64.30%	165.9%
27. N_I	0.0043	−141.66%	177.1%	−155.72%	177.1%
28. $C_I^{(T)}$	0.009	−107.43%	23.0%	−109.47%	17.9%

29. N_{aa}	0.007	-11.19%	21.9%	-12.38%	21.2%
30. C_{ch}	0.0313	-76.26%	161.9%	-45.07%	150.4%
31. C_{su}	0.0313	-75.81%	176.1%	-68.58%	174.1%
32. C_{pr}	0.03	-4.65%	65.7%	-22.21%	69.2%
33. C_{aa}	0.03	-83.80%	178.5%	-95.75%	182.4%
34. C_{li}	0.022	-76.93%	162.2%	-47.79%	163.2%
35. C_{fa}	0.0217	-39.34%	116.6%	-46.52%	123.9%
Kinetic parameters					
36. k_{La}	200	2.12%	8.5%	0.82%	10.3%
37. k_{dis}	0.5	-5.75%	14.4%	-21.53%	26.9%
38. $k_{hyd,ch}$	10	-1.70%	8.5%	-4.78%	14.0%
39. $k_{hyd,pr}$	10	-0.03%	2.8%	-0.56%	4.2%
40. $k_{hyd,li}$	10	0.12%	8.8%	-1.05%	12.8%
41. $k_{m,su}$	30	-103.60%	163.2%	-71.84%	160.1%
42. $K_{S,su}$	0.5	-114.44%	165.8%	-77.91%	153.5%
43. $k_{m,aa}$	50	-90.77%	174.7%	-104.84%	176.9%
44. $K_{S,aa}$	0.3	-86.70%	173.7%	-96.34%	174.4%
45. $k_{m,fa}$	6	-63.78%	93.0%	-77.68%	111.8%
46. $K_{S,fa}$	0.4	-75.58%	129.8%	-104.79%	154.3%
47. $K_{I,h2,fa}$	5.00E-6	-84.41%	175.8%	-141.24%	178.2%
48. $k_{m,c4}$	20	-77.34%	132.9%	-85.21%	153.3%
49. $K_{S,c4}$	0.2	-92.57%	154.2%	-96.66%	163.7%
50. $K_{I,h2,c4}$	1.00E-5	-204.52%	152.8%	-144.35%	178.0%
51. $k_{m,pro}$	13	-93.31%	121.4%	-89.83%	127.8%
52. $K_{S,pro}$	0.1	-112.80%	152.2%	-103.68%	155.7%
53. $K_{I,h2,pro}$	3.50E-6	-116.53%	172.2%	-172.49%	167.5%
54. $k_{m,ac}$	8	-151.33%	115.3%	-150.37%	129.7%
55. $K_{S,ac}$	0.15	-9.02%	36.7%	-17.56%	69.5%
56. $K_{I,nh3}$	0.0018	2.63%	131.4%	-26.54%	151.7%
57. $k_{m,h2}$	35	-109.12%	162.7%	-117.56%	164.6%
58. $K_{S,h2}$	7.00E-6	-77.93%	154.6%	-68.35%	148.5%
59. $k_{dec,Xsu}$	0.02	-24.26%	94.1%	-77.27%	142.6%
60. $k_{dec,Xaa}$	0.02	-29.80%	80.0%	-70.16%	132.0%
61. $k_{dec,Xfa}$	0.02	-11.77%	80.9%	-49.62%	127.2%
62. $k_{dec,Xc4}$	0.02	-2.87%	44.9%	-29.30%	84.2%
63. $k_{dec,Xpro}$	0.02	-7.94%	61.4%	-15.34%	86.6%
64. $k_{dec,Xac}$	0.02	3.66%	50.9%	-24.18%	87.0%
65. $k_{dec,Xh2}$	0.02	9.20%	62.5%	-29.99%	108.7%
66. $pH_{UL,aa}$	5.5	-12.17%	15.2%	-15.85%	15.5%
67. $pH_{LL,aa}$	4	-31.97%	15.3%	-34.53%	16.0%
68. $pH_{UL,h2}$	7	4.59%	14.9%	1.25%	14.5%
69. $pH_{LL,h2}$	6	5.82%	10.7%	2.26%	12.8%
70. $pH_{UL,ac}$	6	-6.92%	16.0%	-14.15%	18.1%
71. $pH_{LL,ac}$	5	-11.53%	12.9%	-13.30%	12.5%
Initial conditions					
72. $S_{su}(0)$	0.00762	0.53%	14.3%	0.64%	27.4%
73. $S_{aa}(0)$	0.0034	-0.97%	15.9%	2.01%	25.3%
74. $S_{fa}(0)$	0.0573	-1.18%	14.1%	-0.08%	20.2%
75. $S_{va}(0)$	0.00725	1.47%	10.8%	0.32%	16.8%
76. $S_{bu}(0)$	0.00826	-0.44%	10.6%	1.32%	15.2%
77. $S_{pro}(0)$	0.00937	-0.31%	10.4%	0.42%	16.7%
78. $S_{ac}(0)$	0.0954	-0.95%	10.8%	-1.71%	20.5%
79. $S_{h2}(0)$	1.500E-7	-0.52%	13.3%	-2.52%	22.3%
80. $S_{ch4}(0)$	0.0505	2.73%	14.6%	5.25%	24.8%

81. $S_{IC}(0)$	0.157	-0.02%	1.4%	1.15%	6.8%
82. $S_{IN}(0)$	0.1337	-0.03%	1.3%	0.84%	6.2%
83. $S_I(0)$	0.422	-0.28%	2.7%	0.36%	7.2%
84. $X_{xc}(0)$	0.201	1.10%	7.3%	2.38%	14.1%
85. $X_{ch}(0)$	0.0145	-1.18%	14.9%	-3.71%	25.9%
86. $X_{pr}(0)$	0.052	-3.29%	13.6%	-3.15%	24.6%
87. $X_{li}(0)$	0.0155	-0.32%	14.8%	-0.90%	26.3%
88. $X_{su}(0)$	0.339	0.30%	2.3%	1.28%	4.0%
89. $X_{aa}(0)$	0.928	0.38%	2.1%	1.29%	4.2%
90. $X_{fa}(0)$	0.2	-0.33%	2.4%	0.70%	5.3%
91. $X_{c4}(0)$	0.341	-0.18%	1.9%	0.36%	5.3%
92. $X_{pro}(0)$	0.109	0.14%	2.8%	0.37%	3.7%
93. $X_{ac}(0)$	0.608	0.21%	2.2%	0.73%	6.6%
94. $X_{h2}(0)$	0.253	-0.55%	2.1%	-0.08%	3.2%
95. $X_I(0)$	25.8	0.00%	2.5%	1.34%	7.4%
96. $S_{cat+}(0)$	0.04	0.02%	2.0%	0.84%	7.1%
97. $S_{an-}(0)$	0.02	0.11%	2.3%	-0.13%	5.1%
98. $S_{gas,h2}(0)$	7.108E-6	-2.14%	15.1%	-0.29%	24.0%
99. $S_{gas,ch4}(0)$	1.582	-1.12%	15.2%	-0.78%	23.5%
100. $S_{gas,co2}(0)$	0.0138	1.44%	13.6%	0.68%	21.3%

Experiment 2

Generally speaking, the results are acceptable. Some interesting results can be obtained for certain subsets. The degradation kinetics of the particulates and the liquid-gas mass transfer coefficient (36.-40.) are estimated well for the noise levels 5%-15%. At 25% these results become very poor for the disintegration constant k_{dis} and $k_{hyd,li}$. Concerning the pH-inhibition levels (65.-71.), estimates for amino acid degradation (66.-67.) and acetoclastic methanogenesis (70.-71.) are accurate and precise for all noise levels. Some variation is observed in the levels for the hydrogenotrophic methanogenesis step (69.-70.). At 25% this variation can be considered to be bad. However, for lower noise levels, the estimates can still be considered to be moderately good. For the sugar and amino acid degradation, the kinetic parameters (41.-44.) as well as the associated stoichiometric parameters (14.-22.) are estimated well for noise levels 5%-15%. At 25%, the estimates are still moderately accurate. The estimates of the other kinetic parameters (45.-65.) are generally poor, regardless of the noise level. The stoichiometric parameters concerning the disintegration of X_{xc} (9.-13.) are generally also poor, regardless of the noise level. The stoichiometric biomass yield parameters have inaccurate estimates, except for the biomass yields of monosaccharides and amino acid degraders (7.-8.) for 5% and 10% noise. The C- and N-composition (23.-35.) estimation results are mixed. The composition of monosaccharide and amino acids (29., 33., 35) is generally well estimated, except for the noise level of 25%. The other compositions show appalling results. Finally, the initial conditions (72.-98.) are moderately well estimated for 5% and 10% noise, except for the acetic acid concentrations (78.) and the LCFA concentrations (74.). For higher noise levels, the estimates generally become unacceptable.

The results can be compared to the covariances of Experiment 1 shown in

Figure 2. There are some similarities: for instance a large part of the kinetic parameters (45.-65.) still shows a great interdependence with each other, i.e., the stoichiometric parameters (9.-13.), the biomass yields (2.-8.) and the composition parameters (23.-35.). There are some differences though. First, the estimates of the initial conditions (72.-98.) now have considerable interdependency with themselves *and* with the other parameters. This is in line with the results in Appendix D that show that the initial conditions estimates are only moderately good. It is also clear that there are more groups of parameters that exhibit little or no interdependency with other parameters. These are the stoichiometric (14.-21.) and kinetic parameters (41.-44.) associated with the degradation of S_{su} and S_{aa} , the first-order kinetic parameters (36.-40.), 4 of the pH-inhibition levels, (66.-67., 70.-71.), and several initial conditions (79.-83., 94.-98.). On the other hand, there are several parameters that show interdependencies with a large set of other parameters. These are the biomass yields (2., 6.) associated with the degradation of S_{ac} and S_{fa} , the composition of X_{xc} (23.-24.), X_{ch} (23.), X_{li} (34.) and S_{fa} (35.). Finally, the pH-inhibition levels of the hydrogenotrophic methanogenesis (68.-69.) show considerable interdependency with the other parameters.

Table 3: Parameter estimation results of Experiment 2 for Gaussian measurement noise with a magnitude of 5% and 10% of the initial conditions. The true parameter values (θ_i^*), the relative deviation of the Monte Carlo estimate $\hat{\theta}_{MC,i\theta}$ and the relative standard deviation of the Monte Carlo estimate $\sigma_{MC,i\theta}$ are shown. $(\dagger)C_I = f_{sI,xc}C_{sI} + f_{xI,xc}C_{xI}$. A bias or standard deviation of less than 5%, between 5% and 25% and more than 25% is indicated in white, light gray and dark gray respectively.

θ_{i_θ}		5%		10%	
	$\theta_{i_\theta}^*$	Bias _{rel,i_θ}	σ_{rel,i_θ}	Bias _{rel,i_θ}	σ_{rel,i_θ}
Stoichiometric parameters					
1. $f_{fa,li}$	0.95	−1.80%	3.6%	−0.41%	4.9%
2. Y_{ac}	0.05	−12.91%	29.5%	−0.45%	34.1%
3. Y_{h2}	0.06	10.64%	2.9%	10.09%	4.7%
4. Y_{pro}	0.04	−0.89%	9.9%	−0.34%	6.5%
5. Y_{c4}	0.06	5.23%	9.4%	6.09%	7.2%
6. Y_{fa}	0.06	−16.62%	67.7%	−14.91%	56.8%
7. Y_{su}	0.1	−2.41%	8.5%	0.13%	8.2%
8. Y_{aa}	0.08	−0.99%	3.1%	−1.29%	6.0%
9. $f_{sI,xc}$	0.1	1.24%	1.9%	1.13%	2.7%
10. $f_{ch,xc}$	0.2	−18.49%	40.2%	−13.66%	44.4%
11. $f_{pr,xc}$	0.2	22.19%	20.1%	18.81%	17.0%
12. $f_{li,xc}$	0.3	16.00%	27.5%	4.53%	36.3%
13. $f_{xI,xc}$	0.2	−28.31%	63.5%	−12.50%	49.4%
14. $f_{bu,su}$	0.13	1.35%	1.8%	1.76%	1.5%
15. $f_{pro,su}$	0.27	−0.72%	1.5%	−0.36%	1.3%
16. $f_{ac,su}$	0.41	−0.57%	3.9%	−1.40%	3.2%
17. $f_{h2,su}$	0.19	1.34%	5.3%	2.33%	4.2%
18. $f_{va,aa}$	0.23	−0.48%	0.7%	−0.42%	0.9%
19. $f_{bu,aa}$	0.26	−1.14%	0.8%	−1.03%	0.8%
20. $f_{pro,aa}$	0.05	1.43%	3.9%	2.80%	4.4%
21. $f_{ac,aa}$	0.4	0.96%	2.4%	0.42%	2.0%
22. $f_{h2,aa}$	0.06	−0.77%	7.6%	0.94%	6.2%
23. C_{xc}	0.02786	−2.28%	59.6%	−17.78%	89.3%
24. N_{xc}	0.00269	2.52%	140.9%	−47.86%	177.1%

25. C_{bac}	0.0313	11.29%	6.7%	11.79%	8.3%
26. N_{bac}	0.0057	2.01%	25.1%	11.06%	37.0%
27. N_{I}	0.0043	-63.27%	152.4%	-37.90%	166.2%
28. $C_{\text{I}}^{(\dagger)}$	0.009	-96.40%	12.3%	-97.80%	9.2%
29. N_{aa}	0.007	-2.39%	7.4%	0.56%	8.9%
30. C_{ch}	0.0313	35.18%	103.8%	36.99%	106.6%
31. C_{su}	0.0313	1.11%	0.8%	1.23%	0.9%
32. C_{pr}	0.03	12.24%	26.6%	11.24%	26.4%
33. C_{aa}	0.03	1.14%	0.9%	1.44%	1.3%
34. C_{li}	0.022	-98.35%	38.0%	-105.12%	34.9%
35. C_{fa}	0.0217	10.96%	38.2%	0.86%	45.3%
Kinetic parameters					
36. k_{La}	200	0.68%	4.2%	1.40%	3.4%
37. k_{dis}	0.5	-1.13%	2.4%	-1.24%	3.2%
38. $k_{\text{hyd,ch}}$	10	-1.99%	4.1%	-1.72%	4.3%
39. $k_{\text{hyd,pr}}$	10	0.96%	1.0%	0.86%	0.8%
40. $k_{\text{hyd,li}}$	10	4.63%	7.1%	2.10%	10.2%
41. $k_{\text{m,su}}$	30	3.32%	1.9%	3.58%	2.4%
42. $K_{\text{S,su}}$	0.5	2.50%	2.7%	3.06%	2.9%
43. $k_{\text{m,aa}}$	50	1.77%	4.7%	3.14%	5.3%
44. $K_{\text{S,aa}}$	0.3	0.18%	0.6%	0.25%	0.8%
45. $k_{\text{m,fa}}$	6	-8.11%	41.9%	-4.04%	41.1%
46. $K_{\text{S,fa}}$	0.4	-28.90%	60.7%	-15.50%	41.0%
47. $K_{\text{I,h2,fa}}$	5.00E-6	-21.51%	88.2%	-35.08%	99.1%
48. $k_{\text{m,c4}}$	20	-5.37%	7.6%	-7.21%	10.8%
49. $K_{\text{S,c4}}$	0.2	-138.06%	84.5%	-165.01%	91.4%
50. $K_{\text{I,h2,c4}}$	1.00E-5	-111.37%	92.9%	-134.54%	98.1%
51. $k_{\text{m,pro}}$	13	-9.91%	8.7%	-9.95%	7.3%
52. $K_{\text{S,pro}}$	0.1	-29.60%	16.1%	-29.90%	17.5%
53. $K_{\text{I,h2,pro}}$	3.50E-6	1.67%	16.7%	6.70%	12.3%
54. $k_{\text{m,ac}}$	8	-229.89%	115.2%	-204.38%	138.9%
55. $K_{\text{S,ac}}$	0.15	45.06%	40.5%	47.01%	39.5%
56. $K_{\text{I,nh3}}$	0.0018	52.14%	88.1%	27.47%	123.8%
57. $k_{\text{m,h2}}$	35	37.61%	4.8%	38.00%	7.5%
58. $K_{\text{S,h2}}$	7.00E-6	16.66%	13.2%	20.38%	14.3%
59. $k_{\text{dec,Xsu}}$	0.02	-6.57%	16.3%	-1.38%	16.4%
60. $k_{\text{dec,Xaa}}$	0.02	0.83%	7.8%	-1.27%	17.6%
61. $k_{\text{dec,Xfa}}$	0.02	-4.73%	42.6%	-15.04%	67.2%
62. $k_{\text{dec,Xc4}}$	0.02	-9.27%	24.7%	-17.06%	27.2%
63. $k_{\text{dec,Xpro}}$	0.02	13.31%	17.2%	13.88%	13.6%
64. $k_{\text{dec,Xac}}$	0.02	-13.11%	50.4%	-21.28%	66.0%
65. $k_{\text{dec,Xh2}}$	0.02	5.22%	3.3%	5.76%	2.6%
66. $\text{pH}_{\text{UL,aa}}$	5.5	-0.02%	0.2%	0.03%	0.2%
67. $\text{pH}_{\text{LL,aa}}$	4	0.06%	0.3%	0.15%	0.3%
68. $\text{pH}_{\text{UL,h2}}$	7	-4.69%	20.7%	-0.76%	18.1%
69. $\text{pH}_{\text{LL,h2}}$	6	-5.61%	23.1%	-0.29%	18.0%
70. $\text{pH}_{\text{UL,ac}}$	6	1.67%	0.3%	1.70%	0.4%
71. $\text{pH}_{\text{LL,ac}}$	5	1.42%	0.2%	1.42%	0.3%
Initial conditions					
72. $S_{\text{su}}(0)$	0.00762	2.36%	11.4%	-0.14%	13.2%
73. $S_{\text{aa}}(0)$	0.0034	2.76%	12.0%	1.54%	14.4%
74. $S_{\text{fa}}(0)$	0.0573	-0.03%	18.0%	2.58%	25.4%
75. $S_{\text{va}}(0)$	0.00725	10.93%	12.5%	12.42%	14.6%
76. $S_{\text{bu}}(0)$	0.00826	7.83%	13.5%	8.75%	22.0%

77. $S_{\text{pro}}(0)$	0.00937	3.75%	12.6%	2.89%	19.3%
78. $S_{\text{ac}}(0)$	0.0954	-14.22%	26.5%	-14.46%	34.9%
79. $S_{\text{h2}}(0)$	1.500E-7	-0.72%	5.2%	-0.02%	10.3%
80. $S_{\text{ch4}}(0)$	0.0505	-0.06%	4.8%	0.68%	11.7%
81. $S_{\text{IC}}(0)$	0.157	-0.81%	1.9%	-0.90%	2.6%
82. $S_{\text{IN}}(0)$	0.1337	0.62%	4.2%	2.00%	6.2%
83. $S_{\text{I}}(0)$	0.422	0.15%	1.6%	0.76%	2.3%
84. $X_{\text{xc}}(0)$	0.201	6.43%	13.6%	8.11%	16.2%
85. $X_{\text{ch}}(0)$	0.0145	2.17%	11.7%	1.06%	11.9%
86. $X_{\text{pr}}(0)$	0.052	4.03%	11.3%	2.16%	12.0%
87. $X_{\text{li}}(0)$	0.0155	3.20%	11.6%	4.10%	13.4%
88. $X_{\text{su}}(0)$	0.339	2.61%	9.7%	-0.31%	10.4%
89. $X_{\text{aa}}(0)$	0.928	2.62%	3.6%	2.34%	3.5%
90. $X_{\text{fa}}(0)$	0.2	2.32%	10.7%	1.25%	7.9%
91. $X_{\text{c4}}(0)$	0.341	-17.28%	14.4%	-22.38%	15.6%
92. $X_{\text{pro}}(0)$	0.109	8.20%	1.2%	8.14%	2.5%
93. $X_{\text{ac}}(0)$	0.608	-1.44%	16.3%	-5.80%	23.4%
94. $X_{\text{h2}}(0)$	0.253	-13.80%	3.2%	-13.00%	4.3%
95. $X_{\text{I}}(0)$	25.8	0.56%	1.3%	0.13%	1.5%
96. $S_{\text{gas,h2}}(0)$	7.108E-6	0.57%	5.1%	0.55%	9.4%
97. $S_{\text{gas,ch4}}(0)$	1.582	0.13%	5.3%	1.12%	9.3%
98. $S_{\text{gas,co2}}(0)$	0.0138	-0.22%	4.4%	0.89%	9.9%

Table 4: Parameter estimation results of Experiment 2 for Gaussian measurement noise with a magnitude of 15% and 25% of the initial conditions. The true parameter values (θ_i^*), the relative deviation of the Monte Carlo estimate $\hat{\theta}_{\text{MC},i_\theta}$ and the relative standard deviation of the Monte Carlo estimate $\sigma_{\text{MC},i_\theta}$ are shown. $(^*)C_I = f_{\text{sI,xc}}C_{\text{sI}} + f_{\text{xI,xc}}C_{\text{xI}}$. A bias or standard deviation of less than 5%, between 5% and 25% and more than 25% is indicated in white, light gray and dark gray respectively.

		15%		25%	
θ_{i_θ}	$\theta_{i_\theta}^*$	Bias _{rel,i_θ}	$\sigma_{\text{rel},i_\theta}$	Bias _{rel,i_θ}	$\sigma_{\text{rel},i_\theta}$
Stoichiometric parameters					
1. $f_{\text{fa,li}}$	0.95	−0.39%	5.7%	2.46%	11.0%
2. Y_{ac}	0.05	−1.74%	41.4%	3.14%	83.0%
3. Y_{h2}	0.06	8.42%	11.0%	1.91%	34.3%
4. Y_{pro}	0.04	−2.05%	12.7%	−7.43%	19.9%
5. Y_{c4}	0.06	5.64%	14.4%	0.34%	31.3%
6. Y_{fa}	0.06	−24.88%	73.0%	−69.05%	120.0%
7. Y_{su}	0.1	1.35%	12.4%	4.77%	19.6%
8. Y_{aa}	0.08	−3.71%	20.3%	−11.64%	32.3%
9. $f_{\text{sI,xc}}$	0.1	2.47%	10.1%	9.69%	21.5%
10. $f_{\text{ch,xc}}$	0.2	−23.11%	46.3%	−26.55%	78.3%
11. $f_{\text{pr,xc}}$	0.2	22.59%	24.2%	27.24%	43.1%
12. $f_{\text{li,xc}}$	0.3	7.47%	34.0%	−5.55%	62.5%
13. $f_{\text{xI,xc}}$	0.2	−11.93%	67.0%	2.79%	103.2%
14. $f_{\text{bu,su}}$	0.13	2.09%	3.1%	2.55%	3.9%
15. $f_{\text{pro,su}}$	0.27	−0.12%	2.1%	0.31%	3.7%
16. $f_{\text{ac,su}}$	0.41	−1.94%	5.6%	−4.30%	9.8%
17. $f_{\text{h2,su}}$	0.19	2.92%	7.6%	7.11%	16.2%
18. $f_{\text{va,aa}}$	0.23	−0.74%	1.9%	−1.65%	4.1%
19. $f_{\text{bu,aa}}$	0.26	−1.27%	1.9%	−1.59%	3.1%

20. $f_{\text{pro,aa}}$	0.05	3.99%	9.6%	10.02%	18.1%
21. $f_{\text{ac,aa}}$	0.4	0.46%	2.8%	-0.49%	5.3%
22. $f_{\text{h2,aa}}$	0.06	1.95%	11.1%	8.12%	21.4%
23. C_{xc}	0.02786	-35.13%	109.4%	-84.74%	147.6%
24. N_{xc}	0.00269	-67.12%	190.4%	-86.86%	192.9%
25. C_{bac}	0.0313	9.53%	34.1%	14.09%	62.0%
26. N_{bac}	0.0057	23.04%	43.9%	35.73%	66.9%
27. N_{I}	0.0043	-58.13%	179.3%	-66.18%	189.7%
28. $C_{\text{I}}^{(*)}$	0.009	-97.43%	13.4%	-98.51%	7.2%
29. N_{aa}	0.007	1.34%	9.8%	2.38%	11.8%
30. C_{ch}	0.0313	28.15%	109.3%	-1.53%	126.8%
31. C_{su}	0.0313	1.27%	2.0%	2.60%	7.1%
32. C_{pr}	0.03	13.74%	26.6%	29.41%	36.8%
33. C_{aa}	0.03	0.97%	7.0%	2.41%	9.7%
34. C_{li}	0.022	-111.14%	59.1%	-135.19%	100.3%
35. C_{fa}	0.0217	-1.58%	61.2%	6.45%	82.0%
Kinetic parameters					
36. $k_{\text{L}a}$	200	1.94%	6.6%	5.78%	15.4%
37. k_{dis}	0.5	-2.62%	12.2%	-25.45%	74.5%
38. $k_{\text{hyd,ch}}$	10	-2.70%	4.8%	-3.39%	10.5%
39. $k_{\text{hyd,pr}}$	10	0.90%	1.2%	2.00%	6.0%
40. $k_{\text{hyd,li}}$	10	-1.02%	25.7%	-11.86%	38.4%
41. $k_{\text{m,su}}$	30	3.67%	9.4%	7.05%	12.0%
42. $K_{\text{S,su}}$	0.5	3.67%	5.3%	7.10%	12.9%
43. $k_{\text{m,aa}}$	50	4.48%	8.7%	8.41%	14.3%
44. $K_{\text{S,aa}}$	0.3	0.32%	1.4%	-0.94%	5.1%
45. $k_{\text{m,fa}}$	6	-0.88%	32.8%	-30.41%	116.5%
46. $K_{\text{S,fa}}$	0.4	-25.82%	59.2%	-42.53%	112.5%
47. $K_{\text{I,h2,fa}}$	5.00E-6	-48.36%	116.4%	-101.71%	153.6%
48. $k_{\text{m,c4}}$	20	-9.05%	34.1%	-17.57%	66.1%
49. $K_{\text{S,c4}}$	0.2	-159.23%	88.3%	-180.39%	98.0%
50. $K_{\text{I,h2,c4}}$	1.00E-5	-129.15%	99.8%	-151.35%	124.5%
51. $k_{\text{m,pro}}$	13	-10.21%	21.4%	-5.61%	17.8%
52. $K_{\text{S,pro}}$	0.1	-33.24%	26.8%	-47.09%	63.6%
53. $K_{\text{I,h2,pro}}$	3.50E-6	3.82%	32.3%	-5.28%	70.3%
54. $k_{\text{m,ac}}$	8	-179.34%	152.5%	-102.65%	169.6%
55. $K_{\text{S,ac}}$	0.15	52.67%	37.2%	37.69%	92.9%
56. $K_{\text{I,nh3}}$	0.0018	11.42%	140.5%	-75.02%	174.2%
57. $k_{\text{m,h2}}$	35	32.05%	45.2%	26.13%	60.1%
58. $K_{\text{S,h2}}$	7.00E-6	21.08%	15.9%	34.41%	24.7%
59. $k_{\text{dec,Xsu}}$	0.02	1.22%	24.2%	4.52%	51.6%
60. $k_{\text{dec,Xaa}}$	0.02	-5.39%	40.2%	-34.04%	85.5%
61. $k_{\text{dec,Xfa}}$	0.02	-20.96%	72.6%	-65.01%	130.7%
62. $k_{\text{dec,Xc4}}$	0.02	-14.78%	30.3%	-28.31%	58.6%
63. $k_{\text{dec,Xpro}}$	0.02	11.95%	19.1%	0.51%	44.0%
64. $k_{\text{dec,Xac}}$	0.02	-35.84%	85.9%	-62.58%	124.2%
65. $k_{\text{dec,Xh2}}$	0.02	6.07%	4.8%	6.39%	13.6%
66. $\text{pH}_{\text{UL,aa}}$	5.5	0.04%	0.6%	0.22%	0.8%
67. $\text{pH}_{\text{LL,aa}}$	4	0.22%	0.6%	0.60%	1.1%
68. $\text{pH}_{\text{UL,h2}}$	7	1.49%	20.8%	7.24%	32.2%
69. $\text{pH}_{\text{LL,h2}}$	6	1.25%	21.2%	7.69%	33.5%
70. $\text{pH}_{\text{UL,ac}}$	6	1.59%	1.1%	1.67%	1.8%
71. $\text{pH}_{\text{LL,ac}}$	5	1.30%	1.1%	1.12%	1.5%
Initial conditions					

72. $S_{su}(0)$	0.00762	2.07%	21.6%	-4.91%	42.9%
73. $S_{aa}(0)$	0.0034	1.21%	22.6%	-3.08%	46.4%
74. $S_{fa}(0)$	0.0573	7.61%	30.0%	6.16%	55.0%
75. $S_{va}(0)$	0.00725	7.98%	24.8%	1.67%	57.1%
76. $S_{bu}(0)$	0.00826	2.40%	34.0%	-1.07%	48.7%
77. $S_{pro}(0)$	0.00937	1.77%	26.0%	-12.10%	52.8%
78. $S_{ac}(0)$	0.0954	-14.79%	42.1%	-20.51%	55.7%
79. $S_{h2}(0)$	1.500E-7	-3.12%	19.0%	-3.31%	30.6%
80. $S_{ch4}(0)$	0.0505	-0.08%	14.8%	4.50%	24.7%
81. $S_{IC}(0)$	0.157	-1.86%	4.9%	-2.92%	9.4%
82. $S_{IN}(0)$	0.1337	3.58%	8.6%	6.90%	10.4%
83. $S_I(0)$	0.422	0.43%	5.6%	-1.07%	12.0%
84. $X_{xc}(0)$	0.201	11.66%	19.6%	17.50%	27.3%
85. $X_{ch}(0)$	0.0145	0.85%	17.9%	1.00%	33.2%
86. $X_{pr}(0)$	0.052	1.66%	22.1%	3.14%	40.0%
87. $X_{li}(0)$	0.0155	3.27%	17.3%	8.83%	31.7%
88. $X_{su}(0)$	0.339	-1.60%	14.9%	-4.32%	22.2%
89. $X_{aa}(0)$	0.928	3.67%	7.0%	-1.18%	26.6%
90. $X_{fa}(0)$	0.2	2.61%	11.8%	6.43%	24.7%
91. $X_{c4}(0)$	0.341	-21.31%	15.8%	-23.23%	21.6%
92. $X_{pro}(0)$	0.109	7.44%	6.3%	10.62%	10.5%
93. $X_{ac}(0)$	0.608	-8.65%	28.8%	-18.29%	34.5%
94. $X_{h2}(0)$	0.253	-11.73%	9.8%	-3.13%	27.4%
95. $X_I(0)$	25.8	0.40%	2.1%	-0.40%	3.8%
96. $S_{gas,h2}(0)$	7.108E-6	1.38%	15.6%	-3.67%	31.1%
97. $S_{gas,ch4}(0)$	1.582	0.36%	15.6%	-1.90%	26.9%
98. $S_{gas,co2}(0)$	0.0138	-0.05%	15.1%	-2.19%	26.7%

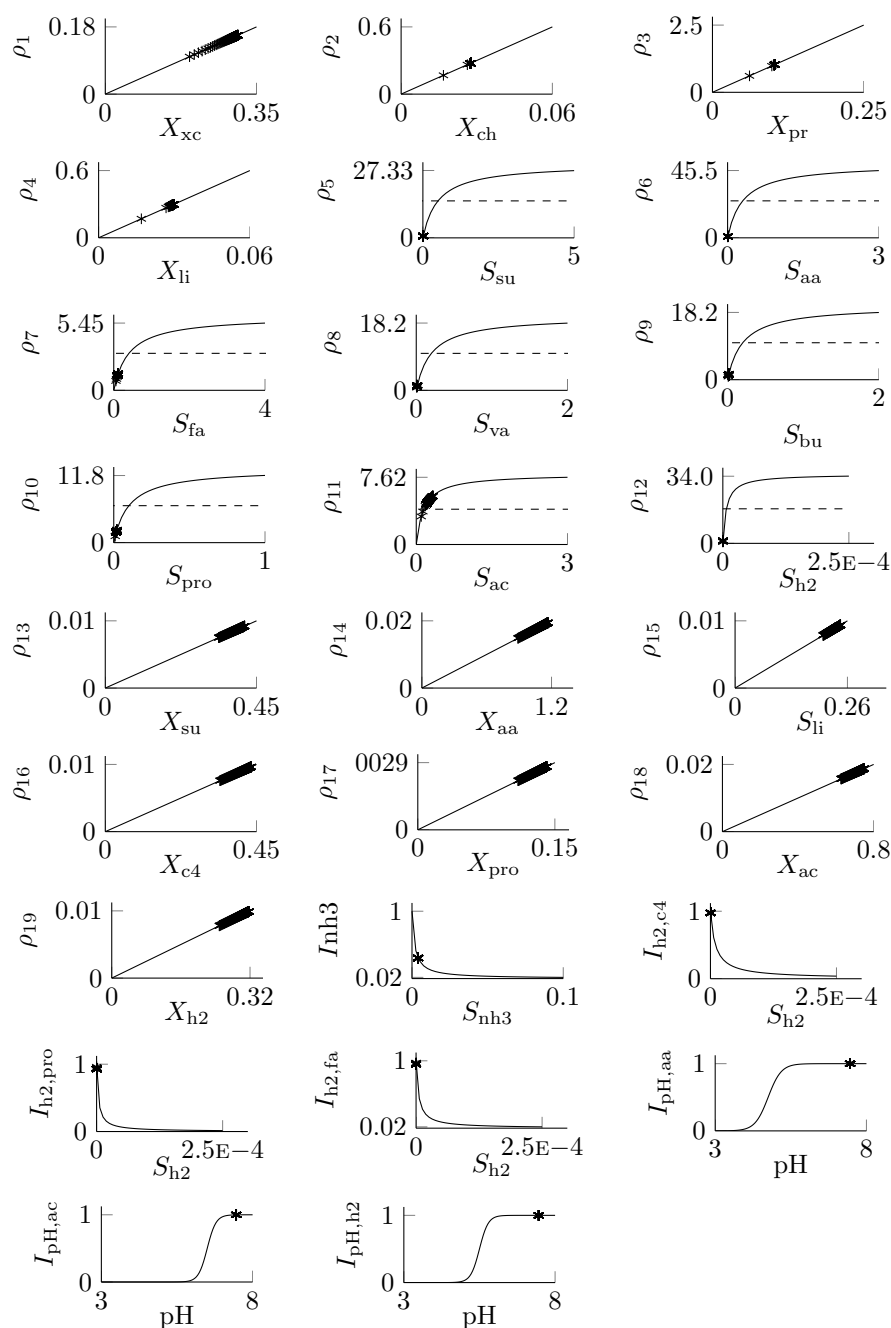


Figure 1: Plot of the simulation results of Experiment 1 on the reaction rates of ADM1. The value $\mu_{max}/2$ is indicated by a dashed line (- -).

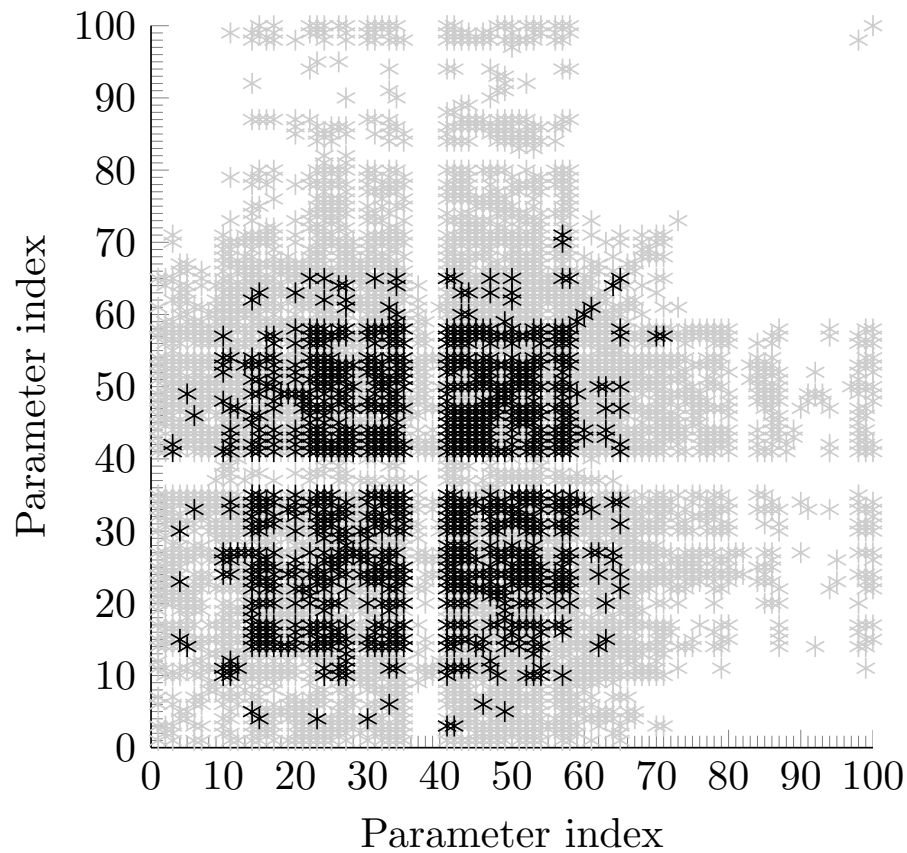


Figure 2: **Square root of the absolute relative covariances for the parameters estimates at a 5% noise level for Experiment 1.** A root of an absolute covariance higher than 25% is indicated in black. A root of an absolute covariance in between 5% and 25% is indicated in grey. Roots of absolute covariances lower than 5 % are not indicated.